L1	148790	S	ASHKENAZI-A?/AU OR BAKER?/AU OR BOTSTEIN-D?/AU OR DESNOYRES?/
Ĺ2	6496	S	GERBER-H?/AU OR GERRITSEN-M?/AU OR GODDARD-A?/AU OR GODOWSKI-
L3	24118	S	KUO-S?/AU OR NAPIER-M?/AU OR PAN-J?/AU OR PAONI-N?/AU OR ROY-
L4	178684	S	L1 OR L2 OR L3
L5	· 2	S	L4 AND PRO363 OR PRO-363
L6	. 2	D	UP REM L5 (0 DUPLICATES REMOVED)
L7 '	16	S	PRO363
L8	0	S	L5 AND 209616
L9	0	S	L4 AND 209616
L10	16	S	PRO363
L11	. 1	S	PRO-363
L12	17	S	L10 OR L11
L13	0	S	L11 AND ANTIBODY

5/4

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Human PRO363 protein sequence SEQ ID NO:503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%;
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GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PAONI N F.
STEPHAN J F.
WATANABE C K.
WILLIAMS P M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GODOWSKI P J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MYERS P S.
KINGSBURY G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BARNES T M.
FRASER C C.
WRIGHTON N.
                                                                 Query Match
Best Local Similarity
RESULT 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             HOLTZMAN D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SHARP J D.
LEIBY K R.
BOSSONE S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MARSTERS S
                                                                                                                                                                                                                                                                                                                                                                                                             protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GURNEY A L
HILLAN K J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GODDARD A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BAKER K
              WO200073454-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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(WILL/)
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(WRIG/)
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(MARS/)
(PANJ/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (PAON/)
(STEP/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (FERR/)
(GERB/)
(GERR/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GODD)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT
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                                                 (without alignments)
1759.288 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Description
                                                                                                              1 MSLLLLLLLVSYYVGTLGTH......TKAETTFSMIPSQSRAFQTV 373
BLOSUM62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          =23-NOV-2000.
(MILL-) MILLENNIUM PHARM INC.
lery Match 100.0%; Score 1908; DB 4; Length 373;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 373;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.0%; Score 1908; DB 3; Length 373; Best Local Similarity 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 373;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 373;
                               Search time 82 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB44248 standard; protein; 373 AA.
Human PRO363 (UNQ318) protein sequence SEQ ID NO:59.
WO200053756-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   / Match 100.0%; Score 1908; DB 3; Local Similarity 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Match 100.0%; Score 1908; DB 4; Local Similarity 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Match 100.0%; Score 1908; DB 2; Local Similarity 100.0%; Pred. No. 3.7e-145.
                                                                                                                                         Gapop 10.0 , Gapext 0.5
Searched: 2105692 seqs, 386760381 residues
Total number of hits satisfying chosen parameters:
Minimum DB seq length: 2000000000
GenCore version 5.1.6
                                                                                                                                                                                                                                   İnimum Match O%
aximum Match 100%
isting first 1500 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB33430 standard; protein; 373 AA.
Human PRO363 protein UNQ318 SEQ ID NO:87.
WO200053758-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY41692 standard; protein; 373 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAU12365 standard; protein; 373 AA. Human PRO363 polypeptide sequence. WO200140466-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein; 373 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAB65293 standard; protein; 373 AA
               OM protein - protein search, using sw m
Run on: May 6, 2005, 10:17:42
                                                                                                                                                                                                                                                                                                                                                                       geneseqp2002s: *
geneseqp2003as: *
geneseqp2003bs: *
                                                                                                                                                                                                                                                                                      A_Geneseq_16Dec04:*
L: geneseqp1980s:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human PRO 363 protein sequence.
WO9946281-A2.
                                                                                                                                                                                                                                                                                                                                         geneseqp2000s:
                                                                                                                                                                                                                                                                                                                                                          geneseqp2001s:
                                                                                US-10-002-967A-59
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-JUN-2001.
(GETH ) GENENTECH INC.
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FOR PAB48108 standard;
                                                                                                                                                                                                                                    Post-processing: M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-SEP-2000
                                                                                                              Sequence:
Scoring table:
                                                                                                  Perfect score:
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Best Local Si
RESULT 2
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                                                                                                                                                                                                                                                                                      Database :
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ID AAY41
DE Humar
PN W0994
PD 16-SI
                                                                               Title:
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Length 373;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABB95454 standard; protein; 373 AA.
Human angiogenesis related protein PRO363 SEQ ID NO:
WO200208284-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABU59186 standard; protein; 373 AA.
Novel human secreted or transmembrane protein PRO363
US2002132252-A1.
O'-DEC-2000.
(GETH ) GENENTECH INC.
(EETH ) GENENTECH INC.
Ery Match 100.0%; Score 1908; DB 4;
                                                                                                                                                          PD 31-JAN-2002.
PA (GETH) GENENTECH INC.
Query Match 100.0%; Score 1908; DB 5;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 8
                                                                                                                                                                                                                                                                                                                                                                                                    100.0%; Score 1908; DB 5; 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                         ABB84848 standard; protein; 373 AA.
Human PRO363 protein sequence SEQ ID NO:64.
WO200200690-A2.
03-JAN-2002.
(GETH ) GENENTECH INC.
                                                                                                                 AAU83656 standard; protein; 373 AA.
Human PRO protein, Seq ID No 130.
WO200208288-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAE26448 standard; protein; 373 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABUSB108 standard; protein; 373 AA.
Human PRO polypeptide #140.
US2003027163-A1.
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Best L RESULT 1

Length 373;

Length 373;

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02-JAN-2003.
(GETH ) GENENTECH INC.
sry Match (GETH ) 100.0%; Score 1908; DB 6; Length 373;
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      100.0%; Score 1908; DB 6; Length 373; 100.0%; Pred. No. 3.7e-145;
                                      Novel human secreted and transmembrane protein PR0363. US2003003531.41. O2-JAN-2003. (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                              ABU84880 standard; protein; 373 AA.
Human secreted and transmembrane polypeptide PRO363.
US2002177553-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     vuery Match
Beet Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 29
                                                                                                                                                               100.0%; Score 1908; DB 6; 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                      100.0%; Score 1908; DB 6; 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABU59844 standard; protein; 373 AA.
Novel secreted and transmembrane protein PRO363.
US2003017563-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABO25034 standard; protein; 373 AA.
Human secreted/transmembrane protein (PRO) #194
US2003036179-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABU80347 standard; protein; 373 AA.
Human secreted/transmembrane protein PRO363.
US2003004102-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABUS9333 standard; protein; 373 AA.
Human secreted/transmembrane protein, #176.
US2003027162-A1.
                                                                                                                                                                                                                      ABUG6763 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003036180-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABO26030 standard; protein; 373 AA.
Human PRO363 polypeptide.
US2002127576-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABUG1078 standard; protein; 373 AA.
Human PRO363 polypeptide.
US2002169284-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-FEB-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                20-FEB-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                         28-NOV-2002.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
        Query Match
Best Local Similarity
RESULT 22
                                                                                                                                                           Query Match
Best Local Similarity
RESULT 23
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                                                                                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 24
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Best Local Similarity
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19-SEP-2002.
(GETH ) GENENTECH INC.
(ery Match 100.0%; Score 1908; DB 6; Length 373;
(ery Match 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       PA (GETH ) GENENTECH INC.
Query Match 100.0%; Score 1908; DB 6; Length 373;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 16
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(GETH ) GENENTECH LTD.
ery Match 100.0%; Score 1908; DB 6; Length 373;
                                                                                                                                                                                 Length 373;
                                                                                                                                                                                                                                                                                                                                             Length 373;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 1908; DB 6; Length 373; 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABU25194 standard; protein; 373 AA.

Novel human secreted and transmembrane protein PRO363.
US2003050239-AI.
13-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel human secreted and transmembrane protein PRO363.
US2003045687-AI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABU72200 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
                                                                                                                                                                                                                                         AB017809 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003032156-A1.
                                                                                                                                                                                    100.0%; Score 1908; DB 6; 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
ery Match
100.0%; Score 1908; DB 6;
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Pred. No. 3.7e-145;
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                                                                                                      ABU82698 standard; protein; 373 AA.
Human secreted/transmembrane protein PRO363.
US2003032023-A1.
13-FEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                  ABUG0617 standard; protein; 373 AA.
Human secreted/transmembrane protein, #176.
US2002160384-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABU81063 standard; protein; 373 AA. Human PRO polypeptide #194. US2003004311-A1. C2-JAN-2003. (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABU80803 standard; protein; 373 AA.
Human PRO polypeptide #65.
US200303635-A1.
20-FBB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein; 373 AA.
                                                                                                                                                                                                                                                                                  ABU13999 standard, protei
Human PRO363 polypeptide.
US2002103125-Al.
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                                                         Best Local Similarity RESULT 13
                                                                                                                                                                                                  Best Local Similarity RESULT 14
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Best Local Similarity
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Best Local Similarity
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BBKER

Length 373;

Query Match

Query Match

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20-MAR-2003.
(GETH ) GENENTECH INC.
bry Match 100.0%; Score 1908; DB 6; Length 373;
st Local Similarity 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                       Length 373;
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                            Length 373;
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Best Local Similarity 100.0%; Pred. No. 3.7e-145;
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                                                                     AUM459U7 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      אם איניבוסאן protein, 373 AA.
Novel human secreted and transmembrane protein PRO363
עוס2003068796-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADA86416 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003082711-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 100.0%; Score 1908; DB 6;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 44
                            100.0%; Score 1908; DB 6; 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                             100.0%; Score 1908; DB 6; 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                       Score 1908; DB 6;
Pred. No. 3.7e-145
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Pred. No. 3.7e-145;
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20-Mar.
                                                                                                                                                                                                                                          ADA76338 standard, protein, 373 AA.
Human PRO polypeptide #194.
US2003073212-A1.
                                                                                                                                                                                                                                                                                                                                                                                                    ABJ72292 standard; protein; 373 AA
Human PRO363 protein.
US2003050448-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADB27937 standard; protein; 373 AA. Human PRO polypeptide #194. US2003082704-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADA61611 standard; protein; 373 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein; 373 AA
                                                                                                                                                                                  100.0%;
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(GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%;
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US2003087350-A1.
                                                                                                                                                                                                                                                                                                      17-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                  (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                  Query Match
Best Local Similarity
RESULT 42
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                                               Best Local Similarity
RESULT 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADB15980 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens.
US2003049816-A1.
          23-JAN-2003.
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                                                                                                                           Length 373;
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RESULT 31

ID ABU82112 standard; protein; 373 AA.

ID ABU82112 standard; protein; 373 AA.

DE Novel human secreted and transmembrane protein PRO363.

PW US2003088063-A1.

PD 08-MAY-2003.

PA (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                      ABU92417 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003022187-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABU59482 standard; protein; 373 AA.
Novel human secreted or transmembrane protein PRO5723
US2003027985-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABUG7039 standard; protein; 373 AA.
Human secreted/transmembrane, PRO, protein SEQ ID 388
US2003032155-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABU92248 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003017476-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABU81706 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2002177164-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABU88645 standard; protein; 373 AA.
Human secreted and transmembrane polypeptide PRO363
US2002197615-A1.
                                                                                                                                                                                                                                            03-OCT-ZUUZ.
(GETH ) GENENTECH INC.
iry Match 100.0%; Score 1908; DB 6;
                                                                            08-MAY-2003.
(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 6;
ery match 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
                                                                                                                                                                            ABUS9039 standard; protein; 373 AA.
Human sectreted/transmembrane protein, #176.
US2002142961-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human PRO polypeptide #140.
05-8PB-70.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             S200302.
13-FEB-2003.
(GETH ) GENENTECH INC.
100.0%;
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28-NOV-2002.
(GETH ) GENENTECH INC.
100.0%;
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PA (GETH) GENENTECH INC.

QUETY MATCh

Best Local Similarity 100.0%;

RESULT 38
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                               Best Local Similarity RESULT 33
                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity RESULT 34
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RESULT 39
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Query Match

RESULT 36

Query Match

Query Match

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Human PRO polypeptide #194.
US2003082763-A1.
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100.0%; Score 1908; DB 6; Length 373;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 59
                                                                                                                                                                                                                                                              UT-APR-2003.
(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 6; Length 373;
       100.0%; Score 1908; DB 6; Length 373; 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Score 1908; DB 6; Length 373; 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADA85864 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003082631-A1.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LUCARRA CONTROL INC. (GETH 100 08; Score 1908; DB 6; sry Match 100 08; Pred. No. 3.7e-145;
                                                                                                              OS-JAN-2003.
(GETH ) GENENTECH INC.
ery Match
100.0%; Score 1908; DB 6;
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                                                                                                                                                                                                                                                                                                                                                       ADA21700 standard; protein; 373 AA.
Human secreted/transmembrane polypeptide PRO363.
US203054404-A1.
20-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADA10487 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
22-WAR-2003.
                                                            ADA38014 standard; protein; 373 AA.
Human secreted/transmembrane protein PRO363
US2003008297-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human PRO polypeptide #194.
US2003082705-Al.
01-MAY-2003.
31TY MATCH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human PRO polypeptide #194.
US2003068795-Al.
                                                                                                                                                                                                           ADA47766 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003073215-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADA18031 standard; protein; 373 AA.
Human PRO353 polypeptide.
US2003054987-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADA79380 standard; protein; 373 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADB30568 standard; protein; 373 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human PRO polypeptide #194 US2003068794-A1.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 53
ID ADA10487 standard; pi
DE Human secreted/transi
PN US2003059831-A1.
PD 27-MRR-2003.
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                  Best Local Similarity
RESULT 50
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RESULT 54
ID ADA
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PN U'
PD PA
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Best Local Similarity 100.0%; Score 1908; DB 6; Length 373; RESULT 68

ID ADB1948 standard; protein; 372 *** DE NOVEL human secrets.
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                                    100.0%; Score 1908; DB 6; Length 373; 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                              y Match 100.0%; Score 1908; DB 6; Length 373; Local Similarity 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADA91813 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003082694-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     _____ADA24598 standard; protein; 373 AA.
ADA24598 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003050241-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADB18837 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363 US2003073211-A1.
                                                                                           ADA87519 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363 US2003087345-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 100.0%; Score 1908; DB 6; Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 1908; DB 6; 100.0%; Pred. No. 3.7e-145;
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(GETH) GENENTECH INC.
(ery Match 100.0%; Score 1908; DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 1908; DB 6; 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145,
                                                                                                                                                                                                                                                                                                                                                                                                          ADA28139 standard; protein; 373 AA.
Human secreted/transmembrane protein PRO363.
US2003054359-A1.
                                                                                                                                                                                                                                        Human PRO polypeptide #194.
08-Mav-and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A. 2514876 standard; protein; 373 AA. Human PRO polypeptide #194.
US2003087351-A1.
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Human PRO polypeptide #194.
US2003077722-Al.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                     08-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 67
                                                      Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 66
                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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Best Local Similarity
01-MAY-2003.
(GETH ) GENENTECH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-MAR-2003.
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                                                                                                                                                                                              Query Match
                                        Query Match
                                                                              RESULT 60
                                                                                                                                                                                                                                                                                                                                                                                        RESULT 62
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Query Match

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01-MXY-2003.
01-MXY-2003.
(GETH ) GENENTECH INC.
ery Match
100.0%; Score 1908; DB 6; Length 373;
                                                                                                                                             Length 373;
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 Length 373;
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                                                                                                                                                                                                                                                                                            Length 373;
                                                    ADAB5312 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003082695-A1.
                                                                                                                                                                                          ADA84760 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363 US2003082708-A1.
 100.0%; Score 1908; DB 6; 100.0%; Pred. No. 3.7e-145;
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Best Local Similarity 100.0%; Pred. No. 3.7e-145;
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(GETH) GENENTECH INC.
(GETH) GENENTECH INC.
ery Match
100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADA38944 standard; protein; 373 AA.
Human secreted/transmembrane protein PRO363.
US2003059780-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADB25307 standard; protein; 373 AA.
Human PRO polypeptide SEQ ID NO 388
US2003077715-A1.
                                                                                                                                                                                                                                                                                                                                  Human PRO polypeptide #194.
17-app.
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US2003082761-A1.
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17-app.
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24-app
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADA75786 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003082703-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                         01-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 83
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                                                                                                                                             Query Match
Best Local Similarity
RESULT 79
               Best Local Similarity RESULT 78
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Best Local Similarity
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   Query Match
                                   Ouery Match 100.0%; Score 1908; DB 6; Length 373; Best Local Similarity 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 6; Length 373;
ery Match 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
rry Match 100.0%; Score 1908; DB 6; Length 373;
rrocal Similarity 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                     ADV#3342 Btandard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003044945-A1.
                                                                                                                                                                                                                                                                                                                                                                                              Novel human secreted and transmembrane protein PRO363 US2003050240-A1.
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20-MAR-2003.
GETH ) GENENTECH INC.
ery Match
100.0%; Score 1908; DB 6;
ery Match
100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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13-MAR-2003.
(GETH) GENENTECH INC.
ery Match
100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA12259 standard; protein; 373 AA.
Human secreted/transmembrane polypeptide PRO363.
US2003055216-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADA94719 standard; protein; 373 AA.
Human secreted/transmembrane protein PRO363
US2003059832-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AUA/5234 standard; protein; 373 AA. Human PRO polypeptide #194. US2003073216-A1. 17-APR-2003. (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADB24747 standard; protein; 373 AA.
Human PRO polypeptide SEQ ID NO 388
US2003077713-A1.
                                                                           Human PRO polypeptide #194.
01-MAV-2000
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Human PRO polypeptide #194.
US2003068798-A1.
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                                                                                                                                                                                                                                                                                                                                                                                     ABO19649 standard; protein; 373 AA
                                                                                                                                                                                                                                                                       US2003v2...
06-MAR-2003.
(GETH ) GENENTECH INC.
""rch 100.0%;
01-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                             01-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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2003.
..tH ) GENENTECH .
.ry Match
Best Local Similarity
RESULT 76
ID ADAS2271 stand-
DE Human PRO r
PN US2003 r
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Best Local Similarity
RESULT 71
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Best Local Similarity
RESULT 72
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Best Local Similarity
RESULT 73
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Best Local Similarity
RESULT 74
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                                                                                        · ADB13260 standard;
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Human PRO polypeptide #194.
US2003082760-A1.
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RESULT
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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 6; Length 373;
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13-MAR-2003.
(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 6; Length 373;
ery Match 100.0%; Pred. No. 3.7e-145;
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ID ADB26281 ct-2-2-2
                                                                   | CETH | GENENTECH INC.
| GETH | GENENTECH INC.
| Ery Match | 100.0%; Score 1908; DB 6; Length 373;
| Core 1908; Dred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 373;
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(GETH) GENERATECH INC.

100.0%; Score 1908; DB 6;

it Local Similarity 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                ADA93065 standard; protein; 373 AA.
Human secreted/transmembrane protein PRO363
US2003060407-A1.
27-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADB24195 standard; protein; 373 AA. Human PRO polypeptide SEQ ID NO 388 US2003077714-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human PRO polypeptide #194.
US2003082690-A1.
                      ADB26833 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003092147-A1.
                                                                                                                                                         ADB31120 standard, protein, 373 AA.
Human PRO polypeptide #194.
US2003096386-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADAB1096 standard, protein, 373 AA Human PRO polypeptide #194. US2003082702-A1. 01-MAY-2003. (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADA95972 standard; protein; 373 AA. Human PRO polypeptide #194.
US2003082759-A1.
01-MAY-2003 (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein; 373 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADB26281 standard; protein; 373 AA
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
RESULT 91
ID ADA61048 standard; p1
DE Homo sapiens.
PN US2003049817-A1.
PD 13-WAR-2003.
PA (GETH ) GENENTECH INC
                                                                                                        Query Match
Best Local Similarity
RESULT 88
ID ADB31120 standard; pl
DE Human PRO polypeptide
PN US2203096586-A1.
PD 22-WAY-2003.
PA (GETH ) GENENTECH INC
                                                                                                                                                                                                                                              Query Match
Best Local Similarity
RESULT 89
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     RESULT 87
ID ADB26
DE Human
PN US200
PD 15-MA
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y Match 100.0%; Score 1908; DB 7; Length 373; Local Similarity 100.0%; Pred. No. 3.7e-145;
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                                      100.0%; Score 1908; DB 6; Length 373; 100.0%; Pred. No. 3.7e-145;
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Novel human secreted and transmembrane protein PRO363.
US2003082700-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADA86968 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003082709-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADA46459 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003054516-A1.
                                                                                               ADB21766 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003082765-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                             ABO19540 standard; protein; 373 AA.
Novel human secreted and transmembrane polypeptide
US2003049633-A1.
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PA (GETH ) GENENTECH INC.

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100.0%; Score 1908; DB 7;

Best Local Similarity 100.0%; Pred. No. 3.7e-145;

RESULT 101
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                                                                                                                                                                                                                                                               ABO34315 standard; protein; 373 AA.
Human secreted/transmembrane polypeptide PRO 363.
US2003044934-A1.
                                                                                                                                                                                                                                                                                                                                                                      Score 1908; DB 6;
Pred. No. 3.7e-145;
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Human PRO polypeptide #194.
US2003077710-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADA77545 standard; protein; 373 AA. Human PRO polypeptide #194. US2003068797-A1.
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Human PRO polypeptide #194.
US2003082699-A1.
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06-MAR-2003.
(GETH ) GENENTECH INC.
100.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 102
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                                                                                                                                                                                                                     Best Local Similarity RESULT 98
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Best Local Similarity
RESULT 99
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Best Local Similarity
RESULT 100
                                                         Best Local Similarity
01-MAY-2003.
(GETH ) GENENTECH
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Score 1908; DB 7; Length 373; Pred. No. 3.7e-145;
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                              Length 373;
                                                                                                                                                                                                                                                                                         100.0%; Score 1908; DB 7; Length 373; 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 1908; DB 7; Length 373;
Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            млависи standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003073213-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel human secreted and transmembrane protein PRO363
US2003087344-A1.
                   Query Match 100.0%; Score 1908; DB 7; Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 1908; DB 7;
Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Human secreted/transmembrane polypeptide PRO363
US2003040473-A1.
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US2003017982-A1.
23-JAN-2003.
                                                                                                                                                                                                                      ABO51245 standard; protein; 373 AA.
Human secreted/transmembrane protein PRO363.
US2003044806-A1.
06-MAR-2003.
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US2003082686-Al.
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US2003022239-A1.
                                                                                 ADB29041 standard; protein; 373 AA. Human PRO polypeptide #194. US2003082706-A1.
                                                                                                                                                                                                                                                                                                                                   Human PRO polypeptide #194. US2003059909-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
              (GETH ) GENENTECH INC.
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Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                      Best Local Similarity
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01-MAY-2003
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ABJ72122 standard; protein; 373 AA.
Human membrane bound receptor/protein PRO363 amino acid sequence.
22003065147-A1.
                                            Length 373;
                                                                                                                                                                      Length 373;
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Novel human secreted and transmembrane protein PRO363 US2003082712-A1.
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US2003073814-A1.
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Novel human secreted and transmembrane protein PRO363
US2003088068-A1.
Human secreted/transmembrane PRO polypeptide #140
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( GETH) GENENTECH INC.

100.0%; Score 1908; DB 7;

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Best Local Similarity 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Query Match
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 119
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
                                                                                                                                                                                                                   ADA39485 standard; protein; 373 AA.
Human secreted/transmembrane protein PRO363.
US2003059782-A1.
                                                                                                                                                                                                                                                                                                                   Human PRO polypeptide #194.
10-2003068793-A1.
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US200307711-Al.
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24-APR-2003.
(GETH ) GENENTECH INC,
100.0%;
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31-MAY-2003.
(GETH ) GENENTECH INC.
100.0%;
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ry Match 100.0%;

r Local Similarity 100.0%;
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Human PRO polypeptide #194.
US2003087352-A1.
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RBSULT 121
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RESULT 117
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                                                          Best Local Similarity RESULT 116
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RESULT 122
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RESULT 124
              US2003049638-A1.
13-MAR-2003.
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Best Local Similarity
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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 7; Length 373;
ery Match 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
100.0%; Score 1908; DB 7; Length 373;
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(GETH ) GENENTECH INC.

100.0%; Score 1908; DB 7; Length 373;

11 Incal Similarity 100.0%; Pred. No. 3.7e-145;
                     Length 373;
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                                                             Novel human secreted and transmembrane protein PRO363.
US2003096968-A1.
                                                                                                                                                                                                                         ADB38680 standard, protein, 373 AA.

Novel human secreted and transmembrane protein PRO363
US2003082766-A1.
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US2003087147-Al.
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Novel human secreted and transmembrane protein PRO363.
US2003082689-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADB78349 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US200309289-A1.
15-MAY-2003.
(GETH) GENENTECH INC.
100.0%; Score 1908; DB 7; Lengt Local Similarity 100.0%; Pred. No. 3.7e-145;
                     Score 1908; DB 7;
Pred. No. 3.7e-145;
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Beet Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 126
                                                                                                                                                                                                                                                                                                                     100.0%; Score 1908; DB 7; 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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US2003054403-A1.
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Human PRO polypeptide #194.
US2003082698-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADB90412 standard; protein; 373 AA. Human PRO polypeptide #194. US2003082762-A1.
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08-MAY-2003.
(GETH ) GENENTECH INC.
100.0%;
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US2003073817-Al.
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(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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                                                                       ID ADB73267 sta
DE Novel human
PN US2003096968
PD · 22-MAY-2003.
PA (GETH ) GENE
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Length 373;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 100.0%; Score 1908; DB 7; Length 373; Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 100.0%; Score 1908; DB 7; Length 373; Best Local Similarity 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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                                              ADB39513 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003082764-A1.
                                                                                                                                                                                                             Novel human secreted and transmembrane protein PRO363.
Novel human secreted and transmembrane protein PRO363.
US2003092886-A1.
15-MAY-2003.
(GETH ) GENENTECH INC.
ery Match
set Local Similarity 100.0%; Pred. No. 3.7e-145;
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Novel human secreted and transmembrane protein PRO363 US2003082687-A1.
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Novel human secreted and transmembrane protein PRO363.
US203069337-A1.
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US200092887-A1.
Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 134
                                                                                                                                                      100.0%; Score 1908; DB 7; 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADB87169 standard; protein; 373 AA.
Human PRO polypeptide #65.
US2003088067-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADB84751 standard; protein; 373 AA.
Human PRO polypeptide #65.
US2003092890-A1.
                                                                                                                                                                                                                                                                                                                                                                                            ADB73565 standard; protein; 373 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB86743 standard; protein; 373 AA. Human PRO polypeptide #194. US2003082697-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                Human PRO polypeptide #8.
US2003045462-A1.
                                                                                                            01-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                        Best Local Similarity
RESULT 135
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ADC57983 standard; protein; 373 AA.
Human PRO polypeptide #140.
US2003027754-A1.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 156
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Best Local Similarity
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RESULT 158
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RESULT 160
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                                               06-FEB-2003.
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                                                      OL-MAY-2003.
(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 7; Length 373;
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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 7; Length 373;
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Pred. No. 3.7e-145;
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                                                                                                                                  ADB77348 standard; protein; 373 AA.
Novel human secreted and transmembrane protein FRO363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel human secreted and transmembrane protein PRO363
US2003082692-A1.
                                                                                                                                                                                                                                                                                                                                     100.0%; Score 1908; DB 7; 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Human secreted/transmembrane protein, PRO363.
US2003054986-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human PRO polypeptide SEQ ID NO 388. US2003077118-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB33953 standard; protein; 373 AA.
Human PRO polypeptide SEQ ID NO 388
US2003077716-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human PRO polypeptide SEQ ID NO 388 24-App.
                                                                                                                                                                                                                                                        ADB34505 standard; protein; 373 AA.
Human PRO polypeptide SEQ ID NO 388
VG2003077717-A1.
                                                                                                                                                                                                                                                                                                                                                                                 ADB35609 standard; protein; 373 AA. Human PRO polypeptide SEQ ID NO 388 US2003077719-A1.
      Human PRO polypeptide #8.
US2003083248-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein; 373 AA
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01-MAZ-2003.
(GETH ) GENENTECH INC.
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20-MAR-2003.
(GETH ) GENENTECH INC.
100.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 145
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Best Local Similarity
RESULT 152
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 RESULT 143
ID ADB762
DE Human
PN US2003
PD 01-MAY
PA, (GETH
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RESULT 147
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RESULT
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Length 373;
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Length 373;
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(GETH) GENENTECH INC.

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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 7;
Score 1908; DB 7;
Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Human secreted/transmembrane protein, PRO363.
US2003054405-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADC66531 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003060406-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jr 159
ADC68655 standard; protein; 373 AA.
Mans secreted/transmembrane protein, PRO363.
US2003064407-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADC62715 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ÀDC67780 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003069178-A1.
                                                                                                                                                                                                                                                                                                                                          Human secreted/transmembrane protein, PRO363
US2003049684-A1.
                                                                                                                                                                                                      Human secreted/transmembrane protein PRO363 US2003049681-A1.
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US2003064375-A1.
                                             Human PRO polypeptide #140.
                                                                                                                                                                                       ADC12214 standard; protein; 373 AA
                                                                                                                                                                                                                                                                                                                       ADC61467 standard; protein; 373 AA
100.0%;
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RESULT 171
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(GETH) GENENTECH INC.
ery Match
ery Match 100.0%; Score 1908; DB 7; Length 373;
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Query Match
100.0%; Score 1908; DB 7; Length 373;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 165
Score 1908; DB 7; Length 373; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                            IT-APR-2003.
(GETH) GENENTECH INC.
2ry Match
2ry Match 100.04; Score 1908; DB 7; Length 373;
2. *^^1 Similarity 100.04; Pred. No. 3.7e-145;
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Novel human secreted and transmembrane protein PRO363.
US2003092106-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC71976 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003092107-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 1908; DB 7;
Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
                                                                                                                                                                                 -DC-1100 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003072745-A1.
                                                                                                                                                                                                                                                                                                                                        ADC67155 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003073131-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC62091 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003073624-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC41724 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003104998-A1.
GETH ) GENENTECH INC.
                                             ADCILG81 standard; protein; 373 AA. Human secreted/transmembrane protein PRO363. US2003069403-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADC21849 standard; protein; 373 AA. Human PRO polypeptide #65.
US2003096969-Al.
22-MAY-2003.
(GETH ) GENENTECH INC.
100.0%; Score 1 bert Local Similarity 100.0%; Pred. N
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Human PRO polypeptide #65.
US2003088065-A1.
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(GETH ) GENENTECH INC.
GETH ) GENENTECH INC.
MATCh ''arity 100.0%;
100.0%;
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(GETH ) GENENTECH INC.
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         Best Local Similarity
RESULT 162
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RESULT 169
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Best Local Similarity
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Query Match
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100.0%; Score 1908; DB 7; Length 373; 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Novel human secreted and transmembrane protein Seg ID388.
US2003087365-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADC57316 standard; protein; 373 AA.
Novel human secreted and transmembrane protein Seg ID388
US2003087366-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ALCOUGES Standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003087361-A1.
ADC59955 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003092105-A1.
IS-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                 ADC49880 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003088064-A1.
                                                                                                                                                                                                                                                                                                                               ADC49079 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003088070-A1.
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Novel human secreted and transmembrane protein PRO363.
US2003088071-A1.
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Novel human secreted and transmembrane protein PRO363
US2003088072-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC60507 standard; protein; 373 AA. Novel human secreted and transmembrane protein PRO363
                                                                                                                                                                                                                             OB-MAY 2003.

(GETH ) GENENTECH INC.
Query Match
BEST Local Similarity 100.0%; Pred. No. 3.7e-145;
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Best Local Similarity 100.0%;
RESULT 180
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                        Best Local Similarity RESULT 172
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Score 1908; DB 7; Length 373; Pred. No. 3.7e-145;
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Novel human secreted and transmembrane protein PRO363
US2002193299-A1.
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Novel human secreted and transmembrane protein PRO363 US2003087354-A1.
                                                                                          ADC9205 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363 US2003087348-A1.
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Novel human secreted and transmembrane protein PRO363
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Pred. No. 3.7e-145;
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US2003083461-A1.
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16-Orm.
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1052003194776-Al.
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Human PRO polypeptide #194.
US2003194770-A1.
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US2003059833-A1.
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) GENENTECH INC.
100.0%; SC
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16-OCT-2003.
(GETH ) GENENTECH INC.
""rch 100.0%; SC
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                             Query Match
Best Local Similarity
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(GETH ) GENENTECH INC.
ry Match 100.0%; Score 1908; DB 7; Length 373;
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                                                                                                                                                   ADC54607 standard; protein; 373 AA.

Novel human secreted and transmembrane protein Seq ID388.
US2003087363-A1.
08-MAY-2003.
(GETH) GENENTECH INC.
100.0%; Score 1908; DB 7; Length
at Local Similarity 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                             ADC59091 standard; protein; 373 AA.
Novel human secreted and transmembrane protein Seq ID388.
US2003087359-A1.
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Novel human secreted and transmembrane protein Seg ID388
US2003087364-A1.
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Novel human secreted and transmembrane protein Seg ID388
US2003087360-A1.
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Novel human secreted and transmembrane protein Seg ID388
US2003087346-A1.
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Novel human secreted and transmembrane protein PRO363.
US2003092104-A1.
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Novel human secreted and transmembrane protein PRO363
US2003082546-A1.
01-MAY-2003.
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Novel human secreted and transmembrane protein PRO363
US2003105288-Al.
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(GETH ) GENENTECH INC.
iry Match 100.0%; Score 1908; DB 7;
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Pred. No. 3.7e-145;
         ADC65509 standard, protein, 373 AA.
Human PRO polypeptide #194.
US2003087362-A1.
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08-MAY-2003.
(GETH) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 7; Length 373;
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Pred. No. 3.7e-145;
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Novel human secreted and transmembrane protein PRO363.
US2003087358-A1.
08-MAY-2003.
(CETH.) GENENTECH INC.
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                                       ADD06312 standard; protein; 373 AA. Novel human secreted and transmembrane protein PRO363. US2003073816-A1.
                                                                                                                                                                                  ADC80573 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003092103-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADD08873 standard; protein; 373 AA.
MOVOL human secreted and transmembrane protein PRO363
US2003073090-A1.
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Novel human secreted and transmembrane protein PRO363
US2003088066-A1.
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(GETH) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 7;
st Local Similarity 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
iry Match 100.0%; Score 1908; DB 7;
ir incal Similarity 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.

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       100.0%; Pred. No. 3.7e-145
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Human secreted/transmembrane PRO polypeptide #32
US2003105011-A1.
05-ZUN-2003.
                                                                                                                                                                                                                                                                                                                                 ADDI1080 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003194774-Al.
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Human PRO polypeptide #194.
US2003194771-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 205
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RESULT I
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100.0%; Score 1908; DB 7; Length 373; 100.0%; Pred. No. 3.7e-145;
                                                                                                        Length 373;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADD50794 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003105291-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADD51040 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US200310520-A1.
05-JUN-2003.
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Novel human secreted and transmembrane protein PRO363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADD41203 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363 US2003203438-A1.
                                                                                                   Query Match 100.0%; Score 1908; DB 7; Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 209
                                                                                                                                                                                                                                                          Query Match 100.0%; Score 1908; DB 7; Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 100.0%; Score 1908; DB 7;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 1908; DB 7; 100.0%; Pred. No. 3.7e-145;
ADD11313 standard; protein; 373 AA.

Human secreted/transmembrane PRO polypeptide #32.
US200310513-A1.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                 Score 1908; DB 7;
Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 1908; DB 7; 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADD52342 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003194769-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADD55476 standard; protein; 373 AA.
Human PRO polypeptide #140.
                                                                                                                                                             ADD09490 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003194775-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADD53082 standard, protein, 373 AA.
Human PRO polypeptide #194.
US2003194792-A1.
                                                                                                                                                                                                                                                                                                                                ADC83369 standard; protein; 373 AA. Human PRO polypeptide #140. US2003059783-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-JUN-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                   16-OCT-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity RESULT 213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
RESULT 215
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
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Best Local Similarity
RESULT 211
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                                                                                                                                                                                                                                                                                                                                                                                                27-MAR-2003.
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Length 373;

Length 373;

RESULT 219

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PA (GETH ) GENENTECH INC.
Query March
100.0%; Score 1908; DB 7; Length 373;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Score 1908; DB 7; Length 373; 100.0%; Pred. No. 3.7e-145;
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                                                                 ADD51286 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003105289-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel human secreted and transmembrane protein PRO363.
US2003194765-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADE27026 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363 US2003087304-A1.
                                                                                                                                                                                                                                                                                                                                                             22 MAY-2003.
(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 7;
ery Match 100.0%; Pred. No. 3.7e-145;
     100.0%; Score 1908; DB 7; 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
                                                                                                                                                                                                               Score 1908; DB 7;
Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                   Ausayuy3 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human secreted/transmembrane protein PRO363.
US2003068647-A1.
10-APR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human PRO polypeptide #194.
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1052003199057-A1.
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USZO03199055-Al.
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US20031990556-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADE31891 standard; protein; 373 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US2003150CT 2003.
23-OCT-2003.
(GETH ) GENENTECH INC.
""" Ch | 100.0%; Sc | 100.0%; Pt | 100.0%;
                                                                                                                             US200310-2003.

05-JUN-2003.

(GETH ) GENENTECH INC,

100.0%;
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PA (GETH) GENENTECH INC.

QUEYY MATCh 100.0%;

Best Local Similarity 100.0%;

RESULT 236
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
RESULT 235
  Query Match
Best Local Similarity
RESULT 227
                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 229
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Best Local Similarity
RESULT 232
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Best Local Similarity
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Best Local Similarity
SSULT 233
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           JT 230
ADD91418 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-MAY-2003.
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                                                                                                                                                                                                                                                                Length 373;
                                                   Length 373;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 373;
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Novel human secreted and transmembrane protein PRO363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22 MAY-2003.
(GETH ) GENENTECH INC.
100.0%; Score 1908; DB 7;
ery Match 100.0%; Pred. No. 3.7e-145;
                                                   100.0%; Score 1908; DB 7; 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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                                                                                                                           ADD37106 standard; protein; 373 AA.
Human secreted/transmembrane PRO polypeptide #32
US2003105012-A1.
                                                                                                                                                                                                                                                                                                                          Human PRO polypeptide #140.
US2003077594-Al.
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16-0CT-2001
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30-0003203431-A1.
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Human PRO polypeptide #65.
US2003096971-A1.
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19-SRD-2002
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Human PRO polypeptide #194.
US2003203430-A1.
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Best Local Similarity 100.0%;
RESULT 222
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US2003096970-A1.
                                                                                                                                                                                                       05-JUN-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                       Best Local Similarity RESULT 218
                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2003203432-A1.
US2003077593-A1.
24-APR-2003.
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                                                      Query Match
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Length 373;

Length 373;

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Human PRO polypeptide #194.
US2003207417-A1.
                                                                                                                     RESULT 246
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(GETH) GENENTECH INC.
ery Match
atch similarity 100.0%; Score 1908; DB 7; Length 373;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCT-2003.
(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 7; Length 373;
ery Match 100.0%; Pred. No. 3.7e-145;
                                                                                                100.0%; Score 1908; DB 7; Length 373; 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                   Length 373;
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Query Match
100.0%; Score 1908; DB 7; Length 373;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-OCT-2003.
(GETH ) GENENTECH INC.
lery Match 100.0%; Score 1908; DB 7; Length 373;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel human secreted and transmembrane protein PRO363. US2001194791-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADE31433 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363 US2003194767-A1.
                                                                                                                                                                                                                                                 Score 1908; DB 7;
Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
ixy Match 100.0%; Score 1908; DB 7;
ixy Match 100.0%; Pred. No. 3.7e-145;
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Best Local Similarity 100.0%; Pred. No. 3.7e-145,
                                                                                                                                     Human secreted/transmembrane protein, PRO363.
192003203434-Al.
                                                                                                                                                                                                                                                                                                    ADE16261 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003203435-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 239

ID ADD72876 standard; protein; 373 AA.

DE Human secreted/transmembrane protein, PRO363.

PN US2003203436-A1.
ADD79485 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003203428-A1.
30-OCT-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADE42021 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003194772-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADE17838 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003199023-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADD91970 standard; protein; 373 AA. Human PRO polypeptide #194.
US2003199053-A1.
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30-CCT-2003.
(GETH ) GENENTECH INC.
CAT MATCh :-:1arity 100.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                         Best Local Similarity
RESULT 237
ID ADE35147 standard; px
DE Human secreted/transm
PN US2003203434-A1.
PD 30-OCT-2003.
PA (GETH ) GENENTECH INC
                                                                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 238
                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
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Best Local Si
RESULT 241
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Best Local Si
RESULT 242
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RESULT 240
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RESULT 245
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100.0%; Score 1908; DB 7; Length 373; 100.0%; Pred. No. 3.7e-145;
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PA (GETH ) GENENTECH INC.
Query Match 100.0%; Score 1908; DB 7; Length 373;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 249
                                                                                                                                                                                                      Query Match 100.0%; Score 1908; DB 7; Length 373; Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 100.0%; Score 1908; DB 7; Length 373; Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Score 1908; DB 7; Length 373; 100.0%; Pred. No. 3.7e-145;
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Novel human secreted and transmembrane protein PRO363.
                                                                                                                                                                                                                                                                                                                                   PD 16-OCT-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 248
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Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 254
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                                                                                                                                                                                                                                                                                                            Human secreted/transmembrane protein, PRO363.
US2003194781-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE18942 standard, protein; 373 AA.
Human PRO polypeptide #194.
US2003199026-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADE43138 standard, protein, 373 AA.
Human PRO polypeptide #194.
US2003199033-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADD95927 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003199059-A1.
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Human PRO polypeptide #194.
US2003199064-A1.
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                                                                                                          ADD93074 standard; protein; 373 AA. Human PRO polypeptide #194. US2003194768-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADE19494 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003199025-A1.
                                                                                                                                                                                                                                                                                       ADD72234 standard; protein; 373 AA
                                                                                                                                                                          16-OCT-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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06-NOV-2003.
(GETH ) GENENTECH INC
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Query Match
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Best Local Similarity
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Best Local Similarity
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Length 373;

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PD .06-NOV-2003.
PA (GETH ) GENENTECH INC.
Query Match
100.0%; Score 1908; DB 7; Length 373;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 267
                                                                                                                                                                                                                                                                                                                                     100.0%; Score 1908; DB 7; Length 373; 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 1908; DB 7; Length 373; 100.0%; Pred. No. 3.7e-145;
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Novel human secreted and transmembrane protein PRO363.
US2003207381-A1.
(GEVVV-2003)
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                    ADG21538 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003207355-A1.
                                                                                                                                                                                                                                                                                                                                                                            Novel human secreted and transmembrane protein PRO363
US2003207384-A1.
                  100.0%; Score 1908; DB 7; 100.0%; Pred. No. 3.7e-145;
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Beet Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 265
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(GETH) GENENTECH INC.
(GETH) GENENTECH INC.
ery Match
100.0%; Pred. No. 3.7e-145;
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                                                                            ADF67430 standard; protein; 373 AA.
Human PRO363 amino acid sequence SEQ ID NO:503.
US2002198148-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADG52656 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003216561-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human secreted/transmembrane protein, PRO363.
US2003206915-A1.
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Human PRO polypeptide #194.
US2003207370-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADG80026 standard; protein; 373 AA. Human PRO polypeptide #194. US2003207372-A1. 06-NOV-2003.
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(GETH ) GENENTECH INC.
  (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 270
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Best Local Similarity
RESULT 266
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Best Local Similarity
RESULT 269
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                                     Best Local Similarity
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                      Query Match
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AD89617 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003199028-A1.
23-OCT-2003.
(GETH) GENENTECH INC.
ery Match
100.0%; Score 1908; DB 7; Length 373;
st Local Similarity 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
2TY MATCh 100.0%; Score 1908; DB 7; Length 373;
St Local Similarity 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
ery Match
100.0%; Score 1908; DB 7; Length 373;
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                                         Length 373;
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                                                                                            ADE32881 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003194766-A1.
                                         100.0%; Score 1908; DB 7; 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 7;
ery Match 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                               ADE16885 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003203433-A1.
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Human secreted/transmembrane protein, PRO363.
US2003195333-A1.
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US2003199031-A1.
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16-0cm
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US2003199032-Al.
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Human PRO polypeptide #194.
US2003199034-A1.
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Human PRO polypeptide #194.
US2003207418-A1.
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16-OCT-2003.
(GETH ) GENENTECH INC.
100.0%;
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PA (GETH) GENENTECH INC.

Query Match 100.0%;

Best Local Similarity 100.0%;

RESULT 257
                                                                                                                                                    16-OCT-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 260
ID ADE40901 standard; pr
DE Human PRO polypeptide
PN US2003199031-A1.
PD 23-OCT-2003.
PA (GETH ) GENENTECH INC
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Best Local Similarity
RESULT 258
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Best Local Similarity
                                                         Best Local Similarity
US2003087305-A1
                  08-MAY-2003.
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Length 373;

Length 373;

Length 373;

Best Loc RESULT 262

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RESULT 289
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100.0%; Score 1908; DB 7; Length 373; 100.0%; Pred. No. 3.7e-145;
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Query Match
100.0%; Score 1908; DB 7; Length 373;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 276
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                                           ADH55870 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003207379-A1.
06-NOV-2003.
(GETH ) GENENTECH INC.
ery Match
st Local Similarity 100.0%; Pred. No. 3.7e-145;
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JT 2/6
Novel human secreted and transmembrane protein PRO363.
US2003207385-A1.
06-NOV-2003.
(GETH ) GENENTECH INC.
100.0%; Score 1908; DB 7; Len
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel human secreted and transmembrane protein PRO363. US2003207388-Al.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AUMB1399 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US203207277-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADI65038 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003207386-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADI63537 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003207387-A1.
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(GETH ) GENENTECH INC.
ry Match 100.0%; Score 1908; DB 7;
....1 cimilarity 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                    Best Local Similarity LOU.UN, LOUST TRESULT 275

RESULT 275

DE Human secreted/transmembrane protein, PRO363.

PN US200377700-A1.

PD 24-APR-2003.
                                                                                                                                                                                  ADI35684 standard; protein; 373 AA.
Human PRO polypeptide #140.
US2003050457-Al.
13-MAR-2003.
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06-NOV-2003.
(GETH ) GENENTECH INC.
100.0%;
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06-NOV-2003.
(GETH ) GENENTECH INC.
"""CA 100.0%;
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(GETH ) GENENTECH INC.
                                                                                                                              Query Match
Best Local Similarity
RESULT 274
ID AD135684 standard, p
DE Human PRO polypeptid
PN US2003050457-A1.
               Best Local Similarity RESULT 273
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Best Local Similarity
RESULT 282
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RESULT I
ID AD:
DE NOV
PN US:
PD 06-
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PD 08-MAY-2003.
AA (GETH ) GENENTECH INC.
Query Match 100.0%; Score 1908; DB 7; Length 373;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 373;
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                                                                                                       100.0%; Score 1908; DB 7; Length 373; 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                 Length 373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 373;
ADM82568 standard; protein; 373 AA.

Novel human secreted and transmembrane protein PRO363.
US2003087355-A1.

(G8-MAY-2003.

(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                          ADNI6596 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003087385-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADNI4863 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003087357-A1.
                                                                                                                                                   Novel human secreted and transmembrane protein PRO363 US2001087353-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADNI5415 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003087356-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADC48833 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003092888-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADC81125 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADE21004 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003100735-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Score 1908; DB 8; 100.0%; Pred. No. 3.7e-145;
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Best Local Similarity 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADD76573 standard; protein; 373 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GETH ) GENENTECH INC.
ry Match
t Local Similarity 100.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                         Best Local Similarity RESULT 283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mest Local Similarity RESULT 286
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US2003100497-A1.
29-MAY-2003.
  US2003100727-A1.
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(GETH ) GENENTECH INC.
ery Match
st Local Similarity 100.0%; Pred. No. 3.7e-145;
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                                                                                  Length 373;
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                                                                                                                                                                                                                                                                                                             ADD75823 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003100717-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADD85055 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADD86881 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
109203100738-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADE20758 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003100734-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADE39055 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003096362-A1.
                                CBTH ) GENENTECH INC.
(GETH ) GENENTECH INC.
ery Match
100.0%; Score 1908; DB 8;
cimilarity 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
ery Match
100.0%; Score 1908; DB 8;
ery match
100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
iry Match 100.0%; Score 1908; DB 8;
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US2003203440-Al.
30-OCT-2003.
(GETH ) GENENTECH INC.
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Human PRO polypeptide #65.
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US2003092113-A1.
                                                                                                                                                protein; 373 AA
Human PRO polypeptide #194
US2003100087-A1.
                                                                                                                                           ADD75077 standard; proteir
Human PRO polypeptide #65.
US2003100712-A1.
                                                                                                                                                                                                           29-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                  Query Match
Best Local Similarity
RESULT 292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
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PD 29-MAY-2003.

PA (GETH ) GENENTECH INC.

Query Match 100.0%; Score 1908; DB 8; Length 373;

Best Local Similarity 100.0%; Pred. No. 3.7e-145;

RESULT 301
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USB (GETH ) GENENTECH INC.

ETY MATCH 100.0%; Score 1908; DB 8; Length 373;

ETY MATCH 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   אבר אים protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003100733-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADD78427 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363 US2003100737-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADE21250 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003100736-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADD77365 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003100732-A1.
                                                                                                                                                                                         Score 1908; DB 8;
Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        PD 05-JUN-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 304
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USCUCED.
(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 8;
ery Match 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADE41314 standard; protein; 373 AA.
Human secreted/transmembrane PRO polypeptide #32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 1908; DB 8;
Pred. No. 3.7e-145
                                                                                                                                                                                                                                                                                                                                                                                                       ADE48393 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003104536-A1.
                                                                             Human PRO polypeptide #65.
US2003100711-A1.
29-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADE23365 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003092108-A1.
                                                                                                                                                                                                                                                      protein; 373 AA
                                                                                                                                                                                           100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%;
                                                                                                                                                                                                                                                                       Human PRO polypeptide #194.
US2003211571-Al.
                                                                                                                                                                           (GETH ) GENENTECH INC.
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                                                                                                                                                                                     Query Match
Best Local Similarity
RESULT 302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
RESULT 309
                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 303
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 306
                                                                                                                                                                                                                                                    ADE75789 standard;
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(GETH ) GENENTECH INC
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                                                                                                                                                                                                                                                                                                   PA (GETH ) GENUTECH INC.

Query Match

Query Match

100.0%; Score 1908; DB 8; Length 373;

Best Local Similarity 100.0%; Pred. No. 3.7e-145;

RESULT 312

ID ADD74339 standard; protein; 373 AA.

DB Human PRO polypeptide #65.

PD 29-MAY-2003.

PA (GETH ) GENENTECH INC.
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ry Match 100.0%; Score 1908; DB 8; Length 373;

t Local Similarity 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                             100.0%; Score 1908; DB 8; Length 373; 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-MAY-2003.
(GETH ) GENENTECH INC.
100.0%; Score 1908; DB 8; Length 373;
it Local Similarity 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AUD76069 standard; protein; 373 AA.

Novel human secreted and transmembrane protein PR0363.
US2003100718-A1.
29-MAY-2003.
(WETH) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADD85561 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003100721-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 100.0%; Score 1908; DB 8; Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 314 D ADD8561 standard; protein; 373 AA. DE Novel human secreted and transmembrane protein PRO: PD 29-MAY-2003.

PA (GETH ) GENEWERFORD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human PRO polypeptide #194.
US2003092110-A1.
                                                                                      ADD75577 standard; protein; 373 AA.
Human PRO polypeptide #65.
US2003100064-A1.
29-MAY-2003.
                                                                                                                                                                                                                       Human PRO polypeptide #65.
US2003100708-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human PRO polypeptide #194.
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Human PRO polypeptide #194.
US2003092111-A1.
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US2003100726-A1.
29-MAY-2003.
                                                                                                                                                             (GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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RESULT 318
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29-MA.
(GETH) ..
Query Match
Best Local $
RESULT 311
ID ADD74r
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PD 7'
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RESULT |
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100.0%; Score 1908; DB 8; Length 373; 100.0%; Pred. No. 3.7e-145;
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Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 373;
Query Match 100.0%; Score 1908; DB 8; Length 373; Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 319
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                                                                                                                                                                                            ADD76867 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003100715-A1.
                                                                                                                                                                                                                                                                                                                             ADD86635 standard, protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003100719-A1.
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                                                                                                                                                                                                                                                                               100.0%; Score 1908; DB 8; 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human PRO polypeptide #194.
US2003199054-A1.
                                                        ADD75323 standard; protein; 373 AA.
Human PRO polypeptide #65.
US2003100714-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADE89251 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003199062-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADE18390 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003194794-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                           29-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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DESNOYERS L.
EATON D L.
FERRARA N.
FILVAROFF E.
FONG S.
                                                                                                                                                       Best Local Similarity RESULT 320
                                                                                                                                                                                                                                                                                            Best Local Similarity RESULT 321
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(BAKE/)
(BOTS/)
(DESN/)
(EATO/)
(FERR/)
(FILV/)
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GIRM/ GURN/ Query Match

TOMA/ SHEL/

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Length 373;
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100.0%; Score 1908; DB 8; Length 373;
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Pred. No. 3.7e-145;
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ry Match 100.0%; Score 1908; DB 8;

t Local Similarity 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
              100.0%; Score 1908; DB 8; 100.0%; Pred. No. 3.7e-145;
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US2003195148-Al.
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Human secreted/transmembrane protein, PRO363
US2003198994-A1.
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23-000.
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                                                            ADE05356 standard; protein; 373 AA.
Human PRO polypeptide #65.
US2003100723-A1.
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Human PRO polypeptide #194.
US2003199027-A1.
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Human PRO polypeptide #194.
US2003199061-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADF35629 standard; protein; 373 AA.
Human PRO363 polypeptide.
US2003194760-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADF45622 standard; protein; 373 AA
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16-OCT-2003
(GETH ) GENENTECH INC.
100.0%; Sr
                                                                                                                                                 100.08;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                 Best Local Similarity RESULT 335
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Best Local Similarity
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                          Best Local Similarity RESULT 334
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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 8; Length 373;
ery Match 100.0%; Pred. No. 3.7e-145;
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US2003100730-A1.
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Novel human secreted and transmembrane protein PRO363.
US2003100725-A1.
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Novel human secreted and transmembrane protein PRO363.
US2003100720-A1.
29-MAY-2003.
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Novel human secreted and transmembrane protein PRO363
US2003100716-A1.
                                                                                                                                                                                                                                                                                                      Score 1908; DB 8;
Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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US2003100713-Al.
29-MAY-2003.
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                                                                                                                                                                                                                                                                                                                                                         ADD77611 standard; protein; 373 AA
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29-MAY-2003.
(GETH ) GENENTECH INC.
100.0%;
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29-MAY-2003.
(GETH ) GENENTECH INC.
""rech 100.0%;
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US2003100710-A1.
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                                                                                                                                                                 PAN J.
PAONI N F.
ROY M A.
STELTON D L.
STEWART T A.
TUMAS D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GETH ) GENENTECH INC.
                                                                                                                   CLJAVIN I J
                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                   KUO S S.
NAPIER M A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                 GURNEY A L
HILLAN K J
                                                                                                                                                                                                                                                                                                                       Local Similarity
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RESULT

(GETH)

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RESULT 354
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RESULT 352
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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 8; Length 373;
                                                                                                                      Length 373;
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US2003199051-A1.
                                                                            OS-CCT_2003.
(GETH ) GENENTECH INC.
(GFT ) GENENTECH INC.
(AFT ) GENENTECH INC.
(AFT ) GENENTECH INC.

100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
(ery Match ) 100.0%; Score 1908; DB 8;
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ry Match
t Local Similarity 100.0%; Pred. No. 3.7e-145;
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Query Match
100.0%; Score 1908; DB 8;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 348
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Best Local Similarity 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
                                                                                                                                                                      ADF24018 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003204055-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF23394 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003203402-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADF33377 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003194780-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human secreted/transmembrane protein, PRO363. US2003199436-A1.
                                                                                                                                                                                                                                                                                                                 ADF40450 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003199021-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADF27480 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003199437-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human PRO polypeptide #194.
US2003199029-Al.
                               ADE93381 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003199060-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /S2003-0-2
30-0CT-2003.
(GETH ) GENENTECH INC.
100.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                        Query Match
Best Local Similarity
RESULT 344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity RESULT 350
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Best Local Similarity
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              RESULT 343
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23-0CT-2003.
(GETH ) GENENTECH INC.
lery Match 100.0%; Score 1908; DB 8; Length 373;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 1908; DB 8; Length 373; 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 1908; DB 8; Length 373; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 1908; DB 8; Length 373; 100.0%; Pred. No. 3.7e-145;
                                                                                                     Length 373;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADE91725 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003199058-A1.
                                                 USZUCZ--
23-CCT-2003.
(GETH ) GENENTECH INC.
(GETY MATCh 100.0%; Score 1908; DB 8;
(ery Match 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%; Score 1908; DB 8;
100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF26220 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003199674-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADF34009 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003194410-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADF46246 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003195344-A1.
                                                                                                                                                                                       Human secreted/transmembrane protein, PRO363
US2003199435-A1.
                                                                                                                                                                                                                                                                                                                                                       Human secreted/transmembrane protein, PRO363 US2003211091-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human secreted/transmembrane protein, PRO363 US2003211092-A1.
ADE90578 standard, protein, 373 AA.
Human PRO polypeptide #194.
US2003199063-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADG11879 standard; protein; 373 AA
                                                                                                                                                                  ADF41074 standard; protein; 373 AA
                                                                                                                                                                                                                                                                                                                                      protein; 373 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein; 373 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PD 11-DEC-2003.
PA (GETH ) GENENTECH INC.
QUERY MALCh
Best Local Similarity 100.0%;
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US2003228655-A1.
                                                                                                                                                                                                                                                                                                                                                                                                  13-NOV-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                             Best Local Similarity
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                                                                                                     Query Match
Best Local Similarity
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ID ADF25119 standard,
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                                                                                                                                                                                                                                                                                                                                    ADF32753 standard;
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Human PRO polypeptide #194.
US2003207353-A1.
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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 8; Length 373;
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(GETH) GENENTECH INC.
ery Match
st Local Similarity 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
iry Match
ir foral Similarity 100.0%; Pred. No. 3.7e-145;
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                                                                                                              Length 373;
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Local Similarity 100.0%; Pred. No. 3.7e-145;
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Local Similarity 100.0%; Pred. No. 3.7e-145;
AUGUD643 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003096559-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADG24283 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003207426-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADG22090 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003207360-A1.
                                                                C2-MAY-2003.
(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 8;
ery Match 100.0%; Pred. No. 3.7e-145;
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Human PRO polypeptide #194.
US2003207352-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADG20160 standard, protein, 373 AA.
Human PRO polypeptide #194.
US2003207376-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADF98066 standard, protein, 373 AA.
Human PRO polypeptide #194.
US2003207422-A1.
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US2003207351-A1.
                                                                                                                                                                       protein; 373 AA.
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                                                                                                                                                               ADG27197 standard; proteir
Human PRO polypeptide #65.
US2003096962-Al.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 363
                                                                                                                                 Best Local Similarity
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PD 06-NOV-2003.

PA (GETH ) GENENTECH INC.

Query Match 100.0%; Score 1908; DB 8; Length 373;

Best Local Similarity 100.0%; Pred. No. 3.7e-145;

RESULT 372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 373;
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                                          Length 373;
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Novel human secreted and transmembrane protein PRO363 US2003096967-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADGO8394 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003207424-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADG12039 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003096963-A1.
                                                                                                                                                                                                                                                                                                                                                     PA (GETH ) GENENTECH INC.
Query Match
Beet Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 373
                                          100.0%; Score 1908; DB 8; 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                               ADG05233 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003207375-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADG13337 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003207357-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADF96962 standard; protein; 373 AA.
Human PRO polypeptide #194.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADG19500 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003207425-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADG15564 standard, protein; 373 AA.
Human PRO polypeptide #194.
US2003219885-A1.
                                                                                                           protein; 373 AA
                                                                                                                             Human PRO polypeptide #194
US2003207359-A1.
06-NOV-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
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Best Local Similarity
                                                          Best Local Similarity RESULT 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity RESULT 375
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ADG60929 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003207309-A1.
06-NOV-2003.
                        Query Match 100.0%; Score 1908; DB 8; Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 389
                                                                                                                                                                                                                                                                                                                                                                    Human PRO polypeptide #194.
US2003207358-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
QUETY MATCh 100.0%;
Best Local Similarity 100.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 391
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity RESULT 395
                                                                                                                                                                                                 Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US2003207362-A1.
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 16-OCT-2003
                                                                                                                                                                                                                                                                                             06-NOV-2003
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                                                                                                                                                                                                   Length 373;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADG07770 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003207356-AI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADG24921 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003207427-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADF94596 standard, protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003096964-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADG07218 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003207350-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADG55265 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003194778-A1.
                                                                                                                                                                                                                                                  ADG23731 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003207389-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vuery Match 100.0%; Score 1908; DB 8; Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 388
                                                    100.0%; Score 1908; DB 8; 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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(GETH) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 8;
ery Match 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
PRY MATCH 100.0%; Score 1908; DB 8;
tt Local Similarity 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.

(GETH ) GENENTECH INC.

(GETH ) GENENTECH INC. 100.0%; Score 1908; DB 8;

(GETH ) GENENTECH INC. 3.7e-145;
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(GETH ) GENENTECH INC.
PY MALCH 100.0%; Score 1908; DB 8;
St Local Similarity 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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US2003207423-A1.
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                                                                                                          protein; 373 AA
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Local Similarity 100.0%;
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US2003207374-A1.
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(GETH ) GENENTECH INC.
                06-NOV-2003.
(GETH ) GENENTECH INC.
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Query Match
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(GETH ) GENENTECH INC.
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RESULT 383
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Best Local Similarity
                                                                   Best Local Similarity RESULT 380
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                                                                                                          ADG06147 standard;
US2003207371-A1.
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Best Local Si
RESULT 381
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100.0%; Score 1908; DB 8; Length 373; 100.0%; Pred. No. 3.7e-145;
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Length 373;
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Pred. No. 3.7e-145;
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Novel human secreted and transmembrane protein PRO363.
US200309665-A1.
22-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADG56921 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003207364-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ALOSSELT standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2002207365-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADG58577 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
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Novel human secreted and transmembrane protein PRO363.
US2003207420-A1.
                                                                   ADG62033 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003207428-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADG57473 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Score 1908; DB 8; 100.0%; Pred. No. 3.7e-145;
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100.0%; Score 1908; DB 8; 100.0%; Pred. No. 3.7e-145;
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Length 373;

Length 373;

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Score 1908; DB 8; Length 373; Pred. No. 3.7e-145;
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                                                                                                                                                                                            ADG52433 standard; protein; 373 AA.

Novel human secreted and transmembrane protein PRO363
US2003207414-A1.
                                                                                                                                                                                                                                                                                                                                                   ADG54161 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363 US2003207416-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADG56369 standard; protein; 373 AA.

Novel human secreted and transmembrane protein PRO363
US2003207366-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADH12635 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003207378-A1.
                                                                                                   20-NOV-2003.
(GETH ) GENENTECH INC.
iry Match
iry Match 100.0%; Score 1908; DB 8;
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  Best Local Similarity 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
                                      ADG51480 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363
US2003215908-A1.
20-NOV-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADG48984 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003216305-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADG48360 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2003216560-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human secreted/transmembrane protein PRO363.
US2003224358-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human PRO polypeptide #194.
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06-NOV-2003.
(GETH ) GENENTECH INC.
"arch 100.0%; SC
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 414
                                                                                                                                                  Best Local Similarity
RESULT 408
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Best Local Similarity
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RESULT 416
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .ADH21242 standard;
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(GETH) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 8; Length 373;
-- rocal Similarity 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
100.0%; Score 1908; DB 8; Length 373;
Fr Match
Fr Local Similarity 100.0%; Pred. No. 3.7e-145;
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                      Length 373;
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                                                                                                                                                                                                                 Auvosidog standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003207415-A1.
                                                                         ADG58025 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003207363-A1.
                                                                                                                                                                                                                                                                                                                                                                                                          Novel human secreted and transmembrane protein PRO363 US2003207421-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADH12011 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003207419-A1.
06-NOV-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 1908; DB 8;
Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
                    100.0%; Score 1908; DB 8; 100.0%; Pred. No. 3.7e-145;
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Best Local Similarity 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADG49608 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
20.200215905-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADH19749 standard; protein; 373 AA.
Human secreted/transmembrane protein PRO363.
US2003228656-A1.
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24-app-2-2-2
                                                                                                                                                                                                                                                                                                                                                                                           protein; 373 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein; 373 AA
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06-NOV-2003.
(GETH ) GENENTECH INC.
Match ''arity 100.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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GENENTECH INC.
                                                                                                                                  06-NOV-2003.
(GETH ) GENENTECH INC
                  Query Match
Best Local Similarity
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Best Local Similarity
RESULT 402
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADG50232 standard;
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                                                                                                                                                                                                                                                                                                                                                                                         ADG71495 standard;
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Length 373;

Query Match

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Human secreted/transmembrane protein, PRO363
                    US2004006219-A1.
08-JAN-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-NOV-2003
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                                                                                                                                                                                                                                                                           RESULT 427
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 431
                                                                                                                                                                                                                                                                                                                                                                         30-JAN-2003.
(GETH) GENENTECH INC.
erry Match
incal Similarity 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GB-JAN-2004.
(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 8; Length 373;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 100.0%; Score 1908; DB 8; Length 373; Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 425
ADG61481 standard; protein; 373 AA.

Novel human secreted and transmembrane protein PRO363.

045203207429-A1.

06-NOV-2003.

(GETH) GENENTECH INC.

100.0%; Score 1908; DB 8; Length 373; sty Match

tocal Similarity 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                         Length 373;
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100.0%; Score 1908; DB 8; Length 373; Best Local Similarity 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADGS4713 standard, protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2003207367-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADG59753 standard, protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2003207369-A1.
                                                                                                                                                                                                            GETH ) GENENTECH INC.

(GETH ) GENENTECH INC.

(GY MATCh

100.0%; Score 1908; DB 8;

---1 Similarity 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
ery Match
100.0%; Score 1908; DB 8;
ery Match
100.0%; Pred. No. 3.7e-145;
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177 Match 1908; DB 8;
17 Match Similarity 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADG50856 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2004005312-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human secreted/transmembrane protein, PRO363.
US2004005657-A1.
                                                                                                                                        Human secreted/transmembrane protein PRO363. 27-Mnv:
                                                                                                                                                                                                                                                                                                            ADG62256 standard; protein; 373 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADH43497 standard; protein; 373 AA. Human PRO polypeptide #32.
US2003224984-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-JAN-2004.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 418
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RESULT 419
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                                                                                                          Best Local Similarity
RESULT 417
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ADH25281 standard, protein, 373 AA.
Human neurotrimin homologue related protein sequence SEQ ID NO:59.
B1386931-A1.
04-FRB-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                          OG-NOV-2003.

(GETH) GENENTECH INC.
Query Match
100.0%; Score 1908; DB 8; Length 373;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 373;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Score 1908; DB 8; Length 373; 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 100.0%; Score 1908; DB 8; Length 373; Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 429
                   Length 373;
                                                                                                                                                                                                                                                                                                                                                                                               Length 373;
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Novel human secreted and transmembrane protein PRO363.
US2003207383-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AD129851 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADG09920 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2004009548-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADI15391 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US200320732-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADG09268 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363
US2004009547-A1.
                                                                                                                                                                                                                                                                                                                                                   PD 22-MAY-2003.

PA (GETH ) GENENTECH INC.

Query Match 100.0%; Score 1908; DB 8;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 428
Query Match 100.0%; Score 1908; DB 8; Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 100.0%; Score 1908; DB 8; Best Local Similarity 100.0%; Pred. No. 3.7e-145; RESULT 433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ouery Match 100.0%; Score 1908; DB 8; Best Local Similarity 100.0%; Pred. No. 3.7e-145;
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Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                                                  Human PRO polypeptide #65.
US200309660-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADH69690 standard; protein; 373 AA.
Human PRO polypeptide #65.
US2004019183-A1.
                                                                                      ADI81177 standard; protein; 373 AA.
Human PRO polypeptide #194.
US2003207361-A1.
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Best Local Similarity 100.0%;
RESULT 434
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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PA (GETH ) GENENTECH INC.
Query Match
100.0%; Score 1908; DB 8; Length 373;
Best Local Similarity 100.0%; Pred. No. 3.7e-145;
RESULT 446
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(HOFF) HOFFMANN LA ROCHE & CO AG F.

ery Match 100.0%; Score 1908; DB 8; Length 373;
                                                                                                                                                                       100.0%; Score 1908; DB 8; Length 373; 100.0%; Pred. No. 3.7e-145;
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Novel human secreted and transmembrane protein PRO363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENENTECH INC.
GENENTECH INC.
h
Aimilarity 100.0%; Pred. No. 3.7e-145;
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                                                                              ADL06892 standard; protein; 373 AA.
Human secreted/transmembrane protein, PRO363.
US2004063921-A1.
                                                                                                                                                                                                                                                                                                                                                                   ADN05373 standard; protein; 373 AA. Antipsoriatic protein sequence #858 WO2004028479-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human PRO polypeptide #194.
US2004077064-A1.
                                                                                                                                                                                                               Human PRO polypeptide #194.
US2004058424-A1.
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US2003077659-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AD036720 standard, protein; 373 AA.
Human UKW polypeptide, SEQ ID NO:2.
EP1416279-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAB48146 standard; protein; 373 AA.
Human A236 variant 2 polypeptide.
WO200069885-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB48147 standard; protein; 373 AA.
Human A236 variant 3 polypeptide.
WO200069885-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                        (GETH ) GENENTECH INC.
                                                                                                                                    01-APR-2004.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                              Query Match
Best Local Similarity
RESULT 444
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Best Local Similarity
RESULT 448
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11-MAR-2004
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                                                                                                                                                                                                                                                                                      04-MAR-2004.
04-MAR-2004.
(GETH ) CENENTECH INC.
ery Match 100.0%; Score 1908; DB 8; Length 373;
ery Match 100.0%; Pred. No. 3.7e-145;
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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 1908; DB 8; Length 373;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADJ77494 standard; protein; 373 AA.

Human PRO polypeptide #194.
US2004038336-A1.
26-FEB-2004.
(GETH ) GENENTECH INC.
100.0%; Score 1908; DB 8; Length 373; stry Match
100.0%; Pred. No. 3.7e-145;
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                                                 100.0%; Score 1908; DB 8; Length 373; 100.0%; Pred. No. 3.7e-145;
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                                                                                                                                                                                                                                       ADM27248 standard; protein; 373 AA.
Novel human secreted and transmembrane protein PRO363.
US2004044179-A1.
                                                                                                                                                                                                                                                                                                                                                                      Novel human secreted and transmembrane protein PRO363.
US2004039164-Al.
                                                                                                                  Novel human secreted and transmembrane protein PRO363 US2003207349-A1.
                                                                                                                                        Score 1908; DB 8;
Pred. No. 3.7e-145;
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Human secreted/transmembrane protein, PRO363.
US2004048332-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human PRO polypeptide #65.
US2004044180-Al.
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26-PPB-26-
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Human PRO polypeptide #194.
US2004048333-A1.
                                                                                                    protein; 373 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /S20040502
26-FEB-2004,
(GETH ) GENENTECH INC.
100.0%;
                22-MAY-2003.
(GETH ) GENENTECH INC.
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                                                            Best Local Similarity
RESULT 435
                                                                                                                                                                                                 Best Local Similarity RESULT 436
                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
RESULT 437
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RESULT 441
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                                                                                                    ADI18318 standard;
US2003096961-A1.
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                                                 Query Match
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Length 373;

Length 373;

Length 373;

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PD 12-UNA-2001.
PA (UYTE-) UNIV TECHNOLOGY CORP.
Query Match 24.4%; 6
Best Local Similarity 31.6%; 1
RESULT 466
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                                                                                                                                                                                 35.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-AUG-1998.
(UYNY ) UNIV NEW YORK STATE.
                                                                                                                                                                                                                                                                                         16-MAY-2002.
(BUHL/) BUHLER T.
(GADI/) GADIENT R A.
(KORN/) KORN R.
(MOVV/) MOVVA R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                             Best Local Similarity RESULT 461
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                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
RESULT 463
                                                                                                                                                                                              Local Similarity
                  (HYSE-) HYSEQ INC.
                                                                                                                                                                (HYSE-) HYSEQ INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human CAR.
US6245966-B1.
 20-MAR-2003
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                                                                                                                                           20-MAR-2003
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                                                                                                                                                                                 Query Match
                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 468
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92.2%; Score 1759.5; DB 4; Length 373; 92.5%; Pred. No. 3.5e-133;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-NOV-2000.
(MILL-) MILLENNIUM PHARM INC.
ery Match 92.3%; Score 1760.5; DB 4; Length 373;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB48150 standard; protein; 373 AA.
Mouse A236 variant 3 polypeptide.
WO20006985-A2.
23-NOV-2000.
(MILL-) MILLENNIUM PHARM INC.
817 MAtch
92.3%; Score 1760.5; DB 4; Length 373;
St Local Similarity 92.5%; Pred. No. 2.9e-133;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               vuery Match 92.3%; Score 1760.5; DB 4; Length 373; Beet Local Similarity 92.5%; Pred. No. 2.9e-133; RESULT 459
                                                                                                                                                                                                                                                                                                            92.4%; Score 1763.5; DB 4; Length 373; 92.7%; Pred. No. 1.7e-133;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92.4%; Score 1763.5; DB 5; Length 373; 92.7%; Pred. No. 1.7e-133;
                                                                                                                                                 Query Match 99.8%; Score 1905; DB 4; Length 373; Best Local Similarity 99.7%; Pred. No. 6.5e-145; RESULT 454
                Length 373;
                99.8%; Score 1905; DB 4; 99.7%; Pred. No. 6.5e-145;
                                                          Human A236 variant 1 polypeptide. W0200069885-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mouse A236 variant 1 polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAB48149 standard, protein, 373 AA.
Mouse A236 variant 2 polypeptide.
WO200069885-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB85862 standard; protein; 373 AA.
Murine adipocytes-derived protein.
WO200166720-A1.
                                                                                                                                                                                                                                                                                                                                                             AAE26449 standard; protein; 373 AA.
Mouse A236 protein.
US2002055139-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AB000483 standard; protein; 141 AA. Novel human polypeptide #70. WO2003023013-A2.
                                                                                                                                                                                                                     AAB48126 standard; protein; 373 AA.
                                                                                                                                                                                                                                      Mouse A236 polyperide.
w020068985-A2.
23-NOV-2000.
(MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MILL-) MILLENNIUM PHARM INC.
Query Match 92.3%;
(MILL-) MILLENNIUM PHARM INC.
                                                                                                                            23-NOV-2000. (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                      (HOLT/) HOLTZMAN D A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SHAR) SHARP J D.
(LEIB/) LEIBY K R.
(BOSS/) BOSSONE S.
(PANY/) PAN Y.
(BARN/) BARNES T M.
(FRAS/) FRASER C C.
(WRIG/). WR.CGTON N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 460
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MYER/) MYERS P S. (KING/) KINGSBURY G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (KITA/) KITAMURA T.
(TSUR/) TSURUGA H.
                          Best Local Similarity RESULT 453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
RESULT 457
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
RESULT 458
                                                                                                                                                                                                                                                                                                                              Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-NOV-2000
                                                                                                                                                                                                                                                                                                          Query Match
Best Local Si
RESULT 455
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                    Query Match
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PD 28-AUG-2003.
PA (USSH) US DEPT HEALTH & HUMAN SERVICES.
Query Match
Best Local Similarity 31.6%; Pred. No. 8e-29;
RESULT 469
    Length 141;
                                                                                                                                                                                         Length 139;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW69697 standard; protein; 365 AA.
Human coxsackievirus and Ad2 and Ad5 receptor HCAR protein.
WO9833819-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 365;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 365;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 365;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24.4%; Score 465; DB 6; Length 365; 31.6%; Pred. No. 8e-29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABJ37063 standard; protein; 365 AA.
Human breast cancer / ovarian cancer related protein #39.
WO2003000012-A2.
36.3%; Score 692; DB 6; 1
100.0%; Pred. No. 1.1e-47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LT 466
ABB08040 standard; protein; 365 AA.
ABB08040 sackie-adenovirus receptor (CAR).
WO200229072-A2.
11-ARR-2002.
(NOVS ) NOVARTIS-ERFINDUNGEN VERW GES MBH.
24 4%; Score 465; DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24.4%; Score 465; DB 2; 31.6%; Pred. No. 8e-29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 2;
                                                                                                                                                                                      Score 670; DB 6;
Pred. No. 6.3e-46;
                                                                                                                                                                                                                                                                                                                                                                                                                                            24.7%; Score 471; DB 5; 32.6%; Pred. No. 2.6e-29;
                                                              AB000802 standard; protein; 139 AA.
Polypeptide encoded by novel human contig #53.
WO2003023013-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 24.4%; Score 465; DB 5.
Best Local Similarity 31.6%; Pred. No. 8e-29;
RESULT 467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW57212 standard; protein; 365 AA.
Human coxsackievirus and adenovirus receptor.
WO9811221-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24.4%; Score 465; DB 2
31.6%; Pred. No. 8e-29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    24.4%; Score 465; DB 4
31.6%; Pred. No. 8e-29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ) 19-MAR-1998.
(DAND ) DANA FARBER CANCER INST INC.
Query Match
                                                                                                                                                                                                                                                          ABB83928 standard; protein; 365 AA.
pCAR SEQ ID NO 4.
US2002059654-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAB47270 standard; protein; 365 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB97544 standard; protein; 365 AA. Human CAR wild-type protein. WO2003070915-A2.
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Length 352;

Length 352;

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ABB95562 standard; protein; 352 AA.
Human angiogenesis related protein PRO5723 SEQ ID NO: 280.
WO200208284-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABUS9187 standard; protein; 352 AA.
Novel human secreted or transmembrane protein PRO5723
US2002132252-A1.
19-SEP-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 433; DB 6;
Pred. No. 2.9e-26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22.7%; Score 433; DB 5; 33.8%; Pred. No. 2.9e-26;
AAB65294 standard; protein; 352 AA.
Human PRO5723 protein sequence SEQ ID NO:505.
WO20073454-A1.
O7-DEC-2000.
A (GETH) GENENTECH INC.
OLEY SCORE 433; DB 4;
BEST LOCAL Similarity 33.8%; Pred. No. 2.9e-26;
                                                                                                                                                                                                                                                          Score 433; DB 5;
Pred. No. 2.9e-26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22.7%; Score 433; DB 6; 33.8%; Pred. No. 2.9e-26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22.7%; Score 433; DB 6; 33.8%; Pred. No. 2.9e-26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22.7%; Score 433; DB 6; 33.8%; Pred. No. 2.9e-26;
                                                                                                                                           Human PRO5723 protein sequence SEQ ID NO:280.
WO200200690-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABU82699 standard; protein; 352 AA.
Human secreted/transmembrane protein PRO5723.
US2003032023-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABUG0618 standard; protein; 352 AA.
Human secreted/transmembrane protein, #177.
US2002160384-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABUS8109 standard, protein, 352 AA.
Human PRO polypeptide #141.
US2003027163-A1.
06-FEB--2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABUBÓ846 standard; protein; 352 AA.
Human PRO polypeptide #108.
US2003036635-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22.7%;
                                                                                                                                                                                                                   PD 03-JAN-2002.
PA (GETH) GENENTECH INC.
Query Match 22.7%;
Best Local Similarity 33.8%;
RESULT 479
                                                                                                                                                                                                                                                                                                                                                                                              GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          GERBER H.
GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (STEP/) STEPHAN J F.
(WATA/) WATANABE C K.
(WILL/) WILLIAMS P M.
(WOOD/) WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GURNEY A L.
HILLAN K J.
MARSTERS S A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GODOWSKI P J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best_Local_Similarity RESULT 481
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
RESULT 483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
RESULT 482
                                                                                                                                                                                                                                                                                                                                                                                                                   BAKER K P.
FERRARA N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PAN J.
PAONI N F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GODDARD A
                                                                                                                                                                                                                                                                                                                                                                           31-JAN-2002
(GETH ) GEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                         (FERR/)
(GERB/)
(GERR/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MARS/)
(PANJ/)
(PAON/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GODD/)
                                                                                                                                                                                                                                                                                                                                                                                                                     BAKE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GURN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ) 19-MAR-1998.

( (DAND ) DANA FARBER CANCER INST INC.

Query Match

Query Match

23.8%; Score 453.5; DB 2; Length 376;

Best Local Similarity 32.0%; Pred. No. 7.1e-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          vuery Match 22.7%; Score 433; DB 4; Length 352;
Best Local Similarity 33.8%; Pred. No. 2.9e-26;
RESULT 476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-DEC-2000.
(GETH ) GENENTECH INC.
ery Match 22.7%; Score 433; DB 4; Length 352;
ery Match 33.8%; Pred. No. 2.9e-26;
                                                                                                                 Score 465; DB 7; Length 365;
Pred. No. 8e-29;
                                                                                                                                                                        D ABU12046 standard; protein; 505 AA.

E Human NOV4a CG59871-01 protein SEQ ID 12.

N WO200281625-A2.

N T-OCT-2002.

A (CURA-) CURAGEN CORP.

Query Match

Best Local Similarity 31.7%; Pred. No. 1.3e-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24.1%; Score 459; DB 8; Length 358; 31.8%; Pred. No. 2.4e-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 358;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-NOV-2000.
MILLENNIUM PHARM INC.
23.9%; Score 456; DB 4; Length 86; Pt Local Similarity 100.0%; Pred. No. 5.8e-29;
                                                                                                                                                                                                                                                                                                                                    ADR66297 standard; protein; 358 AA.
Human prostatic carcinoma derived protein SEQ ID 151 #2.
WO2004076614-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Aur./2018 standard; protein; 352 AA.
Human PRO5723 protein encoded by DNA82361 cDNA clone.
WO20116319-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human BEC/LEC-related protein sequence SeqID148. WO2003080640-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24.1%; Score 459; DB 8; 31.8%; Pred. No. 2.4e-28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAWS7213 standard; protein; 376 AA.
Mouse coxsackievirus and adenovirus receptor.
WO9811221-A2.
    ADN95226 standard; protein; 365 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAB50930 standard; protein; 352 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB48123 standard; peptide; 86 AA.
Human A236 immunoglobulin domain.
WO200069885-A2.
                                                         02-OCT-2003.
(LUDW-) LUDWIG INST CANCER RES.
(LICN) LICENTIA LTD.
                                                                                                                     Query Match
Best Local Similarity 31.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-MAR-2001.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human PRO5723 protein.
WO200073452-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                (HINZ/) HINZMANN B.
(DAHL/) DAHL B.
(ROSE/) ROSENTHAL A.
(HERM/) PERMANN K.
(PILA/) PILARSKY C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-SEP-2004.
(HINZ/) HINZWANN B.
(DAHL/) DAHL E.
(ROSE), ROSENTHAL A.
(HERM/) HERMANN K.
(PILA/) PILARSKY C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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Best Local Similarity
RESULT 477
                                                                                                                                                                                                                                                                                                                                                                                                  10-SEP-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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Length 352;

Length 352;

Length 352;

Length 352;

Length 352;

RESULT 494 ID ABU59483 sta DE Novel human PN US2003027985	PD 06-FEB-2003. Query Match Best Local Simi	RESULT 495 ID ABU92249 sta DE Novel human PN US2003017476 DD 23_IRM-5003	Z e Z		Query Match Query Match Best Local Simi RESULT 497	DE Novel human PN US2002177164 PD 28-NOV-2002. PA (GETH) GENE	282	DE Human secret PN US2002197615 PD 26-DEC-2002.	282	1D ABO34160 SCa DE Human PRO572 PN US2003017981	2 2 2	1D ABO 7.233 BLG DB Human PRO572 PN UGS003050448 PD 13-MAR-2003.	200	DE Human secret PN US2003008297 PD 09-4AN-2003.	ST Be	DE Human secret PN US2003054404 PD 20-MAR-2003.	Best Local Simi RESULT 503 ID ADA10489 sta	DE AUMAII SECTEU PN US2003059831 PD 27-MAR-2003.
6; Length 352; 26;	in PRO5723.	6; Length 352;' 26;		6; Length 352; 26;	in PRO5723.	6; Length 352; 26;		6; Length 352; 26;		6; Length 352; 26;		6; Length 352; 26;	in PRO5723.	6; Length 352; -26;		6; Length 352; 26;	dn PRO5723.	6; Length 352; 26;
INC. 22.7%; Score 433; DB 6; 33.8%; Pred. No. 2.9e-26;	AB033812 standard; protein; 352 AA. Novel human serreted and tränsmembrane protein PRO5723 US2003045687-A1.	INC. 22.7%; Score 433; DB 6; 33.8%; Pred. No. 2.9e-26	protein; 352 AA. peptide.	JTD. 22.7%; Score 433; DB 6; 33.8%; Pred. No. 2.9e-26;	17.48 ABU72585 standard; protein; 352 AA. Novel human secreted and transmembrane protein PRO572: US2003003821-A1.	INC. 22.7%; Score 433; DB 6; 33.8%; Pred. No. 2.9e-26;	protein; 352 AA. ein.	INC. 22.7%; Score 433; DB 6; 33.8%; Pred. No. 2.9e-26;	1, 188 ABUS9334 standard, protein; 352 AA. Human secreted/transmembrane protein, #177. US2003027162-A1.	22.7%; Score 433; DB 6; 33.8%; Pred. No. 2.9e-26;	protein, 352 AA. peptide.	INC. 22.7%; Score 433; DB 6; 33.8%; Pred. No. 2.9e-26	ABU82155 standard; protein; 352 AA. Novel human secreted and transmembrane protein PRO572. US2003088603-A1.	2.7%; Score 433; DB 3.8%; Pred. No. 2.9e	.1 434. ABUSS040 standard; protein; 352 AA. Human sectreted/transmembrane protein, #177. US2002142961-Al. 03-0CTT-2002.	INC. 22.7%; Score 433; DB 6; 33.8%; Pred. No. 2.9e-26;	ABU22418 standard; protein; 352 AA. Novel human secreted and transmembrane protein PRO5723 US2003022187-A1.	22.7%; Score 433; DB 6; 33.8%; Pred. No. 2.9e-26;
PA (GETH) GENENTECH INC Query Match Best Local Similarity RESULT 485	ID ABO33812 standard; DE Novel human secrete PN US2003045687-A1.	See 3		PD 01-ANG-2002. PA (GETH) GENENTECH LTD Query Match Best Local Similarity	KESULT 487 ID ABU72585 standard; DE Novel human secreté PN US2003003531-A1.	g g g		PA (GETH) GENENTECH INC. Query Match Best Local Similarity 33.8%;	IDSOLATION STREET STREE	PD 06-FEB-2003. Query Match Best Local Similarity	ID ABO26031 standard; protein; DE Human PRO5723 polypeptide. PN US2002127576-A1.	PA (GETH) GENENTECH INC. Query Match Best Local Similarity 33.8%;	DB ABU82155 standard; DB Novel human secret; PN US2003088063-A1. PD 08-MAY-2003.	PA (GETH) GENENTECH INC. Query Match Best Local Similarity 3	KESULI 4994 ID ABUS9040 standard; pro DE Human sectreted/transm PN US2002142961-A1. PD 03-0CT-2002.	agg.	ID ABU92418 standard; DE Novel human secretor IN USCO3022187-A1.	PD 30-JAN-2003. Query Match Best Local Similarity

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22.7%; Score 433; DB 6; Length 352; nilarity 33.8%; Pred. No. 2.9e-26;
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22.7%; Score 433; DB 6; Length 352;
illarity 33.8%; Pred. No. 2.9e-26;
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22.7%; Score 433; DB 6; Length 352;
illarity 33.8%; Pred. No. 2.9e-26;
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22.7%; Score 433; DB 6; Length 352;
illarity 33.8%; Pred. No. 2.9e-26;
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1 secreted and transmembrane protein PRO5723.
14-A1.
                                                                                                                                    andard; protein; 352 AA.
secreted and transmembrane protein PRO5723.
6-A1.
andard; protein; 352 AA.
secreted or transmembrane protein PRO3301.
5-A1.
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ted and transmembrane polypeptide PRO5723.
5-A1.
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sted/transmembrane polypeptide PRO5723.
94-A1.
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bted/transmembrane protein, PRO5723.
31-A1.
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ted/transmembrane protein PRO5723.
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23 polypeptide.
1-A1.
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ADA06794 standard; protein; 352 AA.
Human secreted/transmembrane PRO polypeptide #141
US2003049638-A1.
   33.8%; Pred. No. 2.9e-26;
 Best Local Similarity
RESULT 514
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   Score 433; DB 6; Length 352;
Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred: No. 2.9e-26;
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Pred. No. 2.9e-26;
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Human secreted/transmembrane polypeptide PRO 5723.
US2003044934-A1.
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Human secreted/transmembrane polypeptide PRO5723.
US2003040473-A1.
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Human secreted/transmembrane protein PRO5723.
US2003059780-A1.
27-MAR-2003.
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Human secreted/transmembrane protein PRO5723.
US2003060407-A1.
                                                                                                                                                                          ADA28141 standard; protein; 352 AA.
Human secreted/transmembrane protein PRO5723.
US2003054359-A1.
20-MAR-2003.
                                                                                                                                                                                                                                                                                           ADA94721 standard; protein; 352 AA. Human secreted/transmembrane protein PRO5723. US2003059832-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABO53246 standard; protein; 352 AA.
Human secreted/transmembrane protein PRO5723.
US2003044806-A1.
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E Human secreted/transmembrane protein PRO5723.

N US203017982-A1.

D 23-ANN-2003.

Query Match
                                                    ADA18033 standard; protein; 352 AA.
Human PRO5723 polypeptide.
US2003054987-A1.
20-MAR-2003.
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Human PRO5723 protein.
US2003027988-A1.
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06-FEB-2003.
(GETH ) GENENTECH INC.
22.7%; St.
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06-MAR-2003.
(GETH) GENENTECH INC.
22.7%; SC
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33.8%;
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Query Match
Best Local Similarity
RESULT 504
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ABJ72165 standard; protein; 352 AA.
Human membrane bound receptor/protein PRO5723 amino acid sequence.
US2003065147-A1.
Length 352;
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Novel human secreted and transmembrane protein PRO5723.
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Novel human secreted and transmembrane protein PRO5723
US2003088068-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AUB/0435 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723
US2003092899-A1.
                                                                                                                                                                                                                                                                                                                                    Aubsi706 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723
US200073814-A1.
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Novel human secreted and transmembrane protein PRO5723
US20031096968-A1.
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(GETH ) GENENTECH INC.
iry Match 22.7%; Score 433; DB 7;
iry Match 33.8%; Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
Score 433; DB 7;
Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Human secreted/transmembrane protein PRO5723.
US2003059782-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADB96513 standard; protein; 352 AA.
Human PRO polypeptide #141.
US2003054403-A1.
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22.7%;
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Best Local Similarity 33.8%;
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US2003073817-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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ADC36945 standard; protein; 352 AA.
Human PRO polypeptide #108.
US2003088065-A1.
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                                                                                                                                                                         Length 352;
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Pred. No. 2.9e-26;
                                                Length 352;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Novel human secreted and transmembrane protein PRO5723.
US2003092887-A1.
                                                                                                                                                                                                                                                                                                                                            ADB83952 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723.
US2003069397-A1.
                                              Score 433; DB 7;
Pred. No. 2.9e-26;
                                                                                                                                                                         Score 433; DB 7;
Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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DE Human secreted/transmembrane protein PRO5723.
PN US20030649403-A1.
PD '10-APR-2003.
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Human secreted/transmembrane protein PRO5723
US2003049681-A1.
13-MAR-2003.
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                                                                                       ADB87255 standard; protein; 352 AA.
Human PRO polypeptide #108.
US2003088067-A1.
                                                                                                                                                                                                                 ADB84837 standard; protein; 352 AA. Human PRC polypeptide #108. US2003092890.A1. IS-MAY-2003. (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADC57985 standard; protein; 352 AA.
Human PRO polypeptide #141.
US2003027754-A1.
US2003052.
15-MAY-2003.
(GETH ) GENENTECH INC.
22.7%;
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15-MAY-2003.
(GETH ) GENENTECH INC.
MATCh '12-16.
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08-MAY-2003.
(GETH ) GENENTECH INC.
22.7%;
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10-ARP-2003.
(GETH ) GENENTECH INC.
Match '12-ity 33.8%;
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33.8%;
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33.8%;
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                                                           Best Local Similarity RESULT 524
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Score 433; DB 7; Length 352; Pred. No. 2.9e-26;
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Novel human secreted and transmembrane protein PRO5723
US2003088064-A1.
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Novel human secreted and transmembrane protein PRO5723 US2003088070-A1.
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Novel human secreted and transmembrane protein PRO5723
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Novel human secreted and transmembrane protein PRO5723 US2003088072-A1.
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Novel human secreted and transmembrane protein PRO5723
US2003082246-A1.
01-MAY-2003.
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Novel human secreted and transmembrane protein PRO5723 US2003105288-A1.
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Novel human secreted and transmembrane protein PRO5723
US2003068623-A1.
                                                                                                                                                                                              Score 433; DB 7;
Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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                                                                                   Human PRO polypeptide #108.
102003096969-A1.
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                                  Best Local Similarity 33.8%;
RESULT 534
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33.8%;
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(GETH) GENENTECH INC.
Query Match 22.7%;
Best Local Similarity 33.8%;
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08-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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RESULT 542
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Score 433; DB 7; Length 352; Pred. No. 2.9e-26;

Length 352;

Length 352;

Length 352;

Length 352;

552 C083371 standard; protein; 352 AA. mman PRO polypeptide #141. -203059593-A1. -MAR-2003.	\'' a	Z a E				protein; insmembrane INC.	Pred.	Query Match Best Local Similarity 33.8%; Score 433. RESULT 559 ID ADD50607 standard; protein; 352 AA. DE Human PRO polypeptide #108. PN US200396971-A1. PD 22-MAY-2003. PA (GETH) GENENTECH INC.	CH KM PA	PA (GETH) GENENTECH INC. Query Match 22.7%; Score 433; Best Local Similarity 33.8%; Pred. No. 2. RESULT 561 ID ADD50361 standard; protein; 352 AA. DE Human PRO polypeptide #108. PN US2003096970-A1.
PD 01-MAY-2003. Query Match Best Local Similarity 33.8%; Pred. No. 2.9e-26; RESULT 543 ID ADD07804 standard; protein, 352 AA. Novel human secreted and transmembrane protein PROS723.	PD 19-DEC-2002. PD 19-DEC-2002. PA (GETH) GENENTECH INC. Query Match 22.7%; Score 433; DB 7; Length 352; Best Local Similarity 33.8%; Pred. No. 2.9e-26; RESULT 544	DC781 ovel S2003 2-MAY	Sur	Juery Math Surr 546 ADD063 Novel 17-APR	Query Match (SENENTECH INC.) Query Match (Act) Boet Local Similarity 33.8%; Pred. No. 2.9e-26; RESULT 547 ID ADDID1569 standard; protein; 352 AA. DE Human secreted/transmembrane PRO polypeptide #140.	2 g g	Score 433; DB 7; Leng Pred. No. 2.9e-26; 352 AA.	DE Novel human secreted and transmembrane protein PRO5723. PN US2003088066-A1. PD 08-MAY-2003. PA (GETH) GENENTECH INC. Query Match Best Local Similarity 33.8%; Pred. No. 2.9e-26; RESULT 550 ID ADD07124 standard: protein: 352 AA.	2de Beg	ID ADD11529 standard; protein; 352 AA. DE Human secreted/transmembrane PRO polypeptide #140. DE US2003105013-A1. PD 05-UUN-2003. PA (GETH) GENENTECH INC. Query Match 22.7%; Score 433; DB 7; Length 352; Best Local Similarity 33.8%; Pred. No. 2.9e-26;

Score 433; DB 7; Length 352; Pred. No. 2.9e-26;

Score 433; DB 7; Length 352; Pred. No. 2.9e-26;

Length 352;

Score 433; DB 7; Length 352; Pred. No. 2.9e-26;

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RESULT 577
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PN US2003092888-A1.

PD 15-MAY-2003.

PA (GETH) GENENTECH INC.

Query Match

22.7%; Score 433; DB 8; Length 352;

RESULT 571

ID ADE21090 prandand.
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Pred. No. 2.9e-26;
                                            Length 352;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADE26495 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723.
US2003087305-A1.
08-MAY-2003.
                                                                                      ADD51372 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723
US2003105289-A1.
05-JUN-2003.
                                                                                                                                                                                                                                                                                                                                     ADE27028 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723
US2003087304-A1.
08-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE94085 standard; protein; 352 AA.
Immune disease treatment/diagnosis related PRO5723.
US2003082199-A1.
(GETH ) GENENTECH INC.
) 22-WAY-2003.

4 (GETH ) GENENTECH INC.

22.7%; Score 433; DB 7;

Query Match

22.7%; Score 433; DB 7;

Query Match

23.3%; Pred. No. 2.9e-26;
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Human PRO5723 amino acid sequence SEQ ID NO:505.
US2002198148-A1.
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Human secreted/transmembrane protein PRO5723.
US2003068647-A1.
10-APR-2003.
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US2003050457-Al.
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05-JUN-2003.
(GETH) GENENTECH INC.
22.7%; Sc
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26-DEC-2002.
(GETH ) GENENTECH INC.
22.7%;
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                                                      Best Local Similarity
RESULT 562
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Best Local Similarity
RESULT 564
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RESULT 565
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RESULT 567
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Score 433; DB 8; Length 352;
Pred. No. 2.9e-26;
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Novel human secreted and transmembrane protein PRO5723 US2003100735-A1.
29-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                     ADD75909 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723.
US2003100717-A1.
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Novel human secreted and transmembrane protein PRO5723
US2003100722-A1.
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Novel human secreted and transmembrane protein PRO5723
US2003096362-A1.
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Novel human secreted and transmembrane protein PRO5723
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Novel human secreted and transmembrane protein PRO5723
US2003100734-A1.
                                                                                                                                                                                                                                 Score 433; DB 8;
Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
                                                                            Score 433; DB 8;
Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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19-Mav-^^^
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US2003100727-Al.
                                                                                                                                 ADE05934 standard, protein, 352 AA. Human PRO polypeptide #108. 22003100728-AI. 29-MAY-2003. (GETH ) GENENTECH INC.
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Human PRO polypeptide #108.
                                                                                                                                                                                                                           Best Local Similarity 33.8%;
RESULT 573
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(GETH ) GENENTECH INC.
Query Match
Best Local Similarity 33.8%;
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                                                                          Query Match 22.7%;
Best Local Similarity 33.8%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Query Match
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 576
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US20031vv._
29-MAY-2003.
(GETH ) GENENTECH INC.
22.7%; S'
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33.8%;
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PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 33.8%;
RESULT 594
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                 (GETH ) GENENTECH INC.
Query Match 22.7%;
Best Local Similarity 33.8%;
                                                                                                                                                                                                                                                                                                                                             Human PRO polypeptide #108
US2003100714-Al.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                  Best Local Similarity RESULT 591
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Best Local Similarity
SSULT 593
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    29-MAY-2003
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                                                                                                                                                          Query Match
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                                                              RESULT 590
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                                              Score 433; DB 8; Length 352;
Pred. No. 2.9e-26;
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(GETH ) GENENTECH INC.
ery Match 22.7%; Score 433; DB 8; Length 352;
st Local Similarity 33.8%; Pred. No. 2.9e-26;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADD77451 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723.
US2003100732-A1.
                                                                                                           Novel human secreted and transmembrane protein PRO5723
US2003100737-A1.
                                                                                                                                                                                                                                                                                                                                  Aucellate standard, protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723
US2003100736-Al.
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Novel human secreted and transmembrane protein PRO5723
US2003100733-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADD76155 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723
US2003100718-A1.
                                                                                                                                                                                                                    ADE41530 standard; protein; 352 AA.
Human secreted/transmembrane PRO polypeptide #140.
US2003100497-A1.
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Human PRO polypeptide #108.
US2003100709-A1.
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                                                                                             protein; 352 AA
                                                                                                                       US2003100.2
29-MAY-2003.
(GETH ) GENENTECH INC.
22.7%; SC
US200310...
29-MAY-2003.
(GETH ) GENENTECH INC.
22.7%; St.
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29-MAY-2003.
(GETH ) GENENTECH INC.
22.7%; S
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29-MAY-2003.
(GETH ) GENENTECH INC.
22.7%; S
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29-MRY-2003.
(GETH ) GENENTECH INC.
MATCh '13-rity 33.8%;
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(GETH ) GENENTECH INC.
MATCh '-19rity 33.8%;
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29-MXY-2003.
(GETH ) GENENTECH INC.
22.7%;
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ry Match 22.7%;
t Local Similarity 33.8%;
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US2003100064-A1.
29-MAY-2003.
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US2003100708-A1.
                                                           Best Local Similarity
RESULT 581
                                                                                                                                                                                    Best Local Similarity RESULT 582
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Best Local Similarity
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                                                                                           ADD78513 standard;
                                                 Query Match
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Query

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Length 352;
Length 352;
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                                 nowobod7 standard; protein; 352 AA.

Novel human secreted and transmembrane protein PRO5723.
29-MAY-2003.
(GETH ) GENENTECH INC.
17 Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADD86721 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723.
US2003100719-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADD85401 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723
US2003100725-A1.
9-MAY-20072
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADD76953 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PROS723
US2003100715-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel human secreted and transmembrane protein PRO5723 US2003100731-A1.
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Novel human secreted and transmembrane protein PRO5723
US2003100729-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADD77943 standard, protein, 352 AA.
Novel human secreted and transmembrane protein PRO5723
US2003100730-A1.
                                                                                                                                                                                                                                                                                                                        Score 433; DB 8;
Pred. No. 2.9e-26;
Score 433; DB 8;
Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
                                                                                                                                                                                                           Human PRO polypeptide #108.
19-mav-29-may-29-may-2003
                                                                                                                                                                                                                                                                                                                                                                                     ADD75409 standard; protein; 352 AA
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Best Local Similarity 33.8%; Pred. No. 2.9e-26; RESULT 608
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DE Novel human secreted and transmembrane protein PR05723.

PN US2003096959-A1.

PD 22-MAY-2003.

PA (GETH ) GENENTECH INC.

Query Match

22.7%; Score 433; DB 8; Length 352;
                                            Length 352;
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                                                                                                                                                                                                                                                                                                                                                                                                                           ADD77199 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723
US2003100716-A1.
                                            Score 433; DB 8;
Pred. No. 2.9e-26;
                                                                                                                                                               Score 433; DB 8;
Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Human PRO polypeptide #108.
US2003100723-A1.
29-MAY-2003.
(GETH ) GENENTECH INC.
22.7%; Score 43
uery Match
est Local Similarity 33.8%; Pred. No.
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29-Mav-2012
                                                                                        protein; 352 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein; 352 AA
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JIT 606

JIT 606

ADG11881 standard; protein,

Human PRO5723 polypeptide.

( US200328655-A1.

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A (GETH ) GENENTECH INC.

A CETH ) GENENTECH INC.
(GETH ) GENENTECH INC. 22.7%; S
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29-MAY-2003.
(GETH ) GENENTECH INC.
22.7%; S.
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29-MRZ-2003.
(GETH ) GENENTECH INC.
22.7%;
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29-MAY-2003.
(GETH ) GENENTECH INC.
"1-ch "22.7%;
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                                                                                                    Human PRO polypeptide #108
US2003100710-A1.
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US2003100713-A1.
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Human PRO5723 polypeptide.
US2003194760-Al.
16-OCT-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                        ID ADD85893 standard, prote DE Novel human secreted and PN US2003100720-A1.
PD . 29-MAY-2003.
PA (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                   29-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                             LOO3.

LH ) GENENTECH LY MATCH LOCAL SIMILARITY - RESULT 600

ID ADD74671 stands PR Human PRO ry PN US20031
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Best Local Similarity
RESULT 602
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Best Local Similarity
RESULT 604
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Best Local Similarity
RESULT 605
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                                                                                      ADD73933 standard;
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                                             Query Match
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Score 433; DB 8; Length 352;
Pred. No. 2.9e-26;
                                                                                      Length 352;
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                                                                                                                                          ADG11346 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723
US2003096967-A1.
                                                                                                                                                                                                                                                                  AUG12125 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723.
US2003096963-A1.
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Novel human secreted and transmembrane protein PROS723.0520090965-A1.
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Novel human secreted and transmembrane protein PRO5723
US2003096964-A1.
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Pred. No. 2.9e-26;
                                                                                                                                                                                                                                                                                                                                            22-MAY-2005.
(GETH ) GENENTECH INC.
22.7%; Score 433; DB 8;
MATCh 22.7%; Score 433; DB 8;
22.7%; Score 433; DB 8;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
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Pred. No. 2.9e-26
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Human secreted/transmembrane protein PRO5723.
US2003228656-A1.
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Human secreted/transmembrane protein PRO5723.
US2003224358-A1.
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Human secreted/transmembrane protein PRO5723.
US2003219856-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human PRO polypeptide #108.
US2003096966-A1.
ADG27283 standard; protein; 352 AA. Human PRO polypeptide #108.
US2003096962-A1.
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22-MAY-2003.
(GETH ) GENENTECH INC.
45FGh 22.7%; St
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33.8%;
                                                   PD 22-MAY-2003.

PA (GETH ) GENENTECH INC.

QUETY MATCH 22.7%;

BEST LOCAL Similarity 33.8%;

RESULT 609
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Best Local Similarity 33.8%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-NOV-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity RESULT 614
                                                                                                                                                                                                                                              Best Local Similarity RESULT 610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
RESULT 616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
RESULT 617
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-DEC-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-MAY-2003
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Length 352

5

5; Length 261;

8; Length 430;

Length 431;

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ADIOGEO1 standard; protein; 431 AA.
Human coxsackie and adenovirus receptor like 1 (CXADRL1) protein.
WO2003104275-A2.
Mouse coxsackievirus and Ad2 and Ad5 receptor MCAR protein.
W09833819-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABR62054 standard; protein; 431 AA.
Human cell adhesion molecule BT-IgSF protein SEQ ID NO:2.
EP1321475-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABU08259 standard; protein; 431 AA.
Human voltage-gated sodium channel beta subunit HIPHUM30
GB2376235-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PD 25-JUN-2003.
PA (MORG) MORINAGA MILK IND CO LTD.
Query Match
Best Local Similarity 30.2%; Pred. No. 2.4e-20;
RESULT 631
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18.9%; Score 360; DB 5; 30.2%; Pred. No. 2.9e-20;
                                                                                                                                                                                                                                                                              19.8%; Score 378.5; DB 5 34.6%; Pred. No. 4.8e-22;
                                                                                                                                                                                                                                                                                                                                                                                                            CYTOS BIOTECHNOLOGY AG.

19.0%; Score 362.5; DB 8
Similarity 29.5%; Pred. No. 1.8e-20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 361; DB 6;
Pred. No. 2.4e-20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18.9%; Score 360; DB 4; 30.2%; Pred. No. 2.9e-20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18.9%; Score 360; DB 7; 30.2%; Pred. No. 2.9e-20;
                                                                            Score 422.5; DB
Pred. No. 2e-25;
                                                                                                                               ABB83927 standard; protein; 261 AA.
C-terminally truncated pCAR SEQ ID NO
US2002059654-A1.
                                                                                                                                                                                                                                                                                                                               ADL67136 standard; protein; 430 AA.
Human B7-H5 protein SEQ ID NO:6.
WO2004022594-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABB10359 standard; protein; 426 AA.
Human cDNA SEQ ID NO: 667.
WO200154474-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABP66946 standard; protein; 426 AA.
Human polypeptide SEQ ID NO 667.
US2002090672-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein; 431 AA
                                OG-AUG-1998. (UXNY) UNIV NEW YORK STATE. (UXNY) at 22.1%; ery Match ....larity 29.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18.9%;
30.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (INCY-) INCYTE GENOMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ONCO-) ONCOTHERAPY SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-DEC-2002.
(GLAX ) GLAXO GROUP LTD.
                                                                                                                                                                                    16-MAY-2002.
BUHL/) BUHLER T.
(GADI/) GADIENT R A.
(KORN/) KORN R.
(MOVV/) MOVVA R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
RESULT 629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
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                                                                                           Best Local Similarity
RESULT 627
                                                                                                                                                                                                                                                                                               Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ROSE/) ROSEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
RESULT 633
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADC42862 standard;
REMAP protein #22.
WO2003027228-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                           18-MAR-2004
                                                                                                                                                                                                                                                                                Query Match
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                                                                               Query Match
                                                                                                                                                                                                                                                                                                                    RESULT 628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 424.5; DB 6; Length 422;
Pred. No. 1.8e-25;
                                                                                               Length 352;
                                                                                                                                                                                                                                                                                                                                                                                           Score 433; DB 8; Length 352;
Pred. No. 2.9e-26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 352;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 433; DB 8; Length 352;
Pred. No. 2.9e-26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 352;
                                                                                                                                                                                                                                              Length 352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADI29937 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723.
US2003096961-A1.
                                                                                                                                                ADG34212 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723
US2004006206-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADM27334 standard; protein; 352 AA.
Novel human secreted and transmembrane protein PRO5723
US20040444179-A1.
                                                                                                                                                                                                                                            Score 433; DB 8;
Pred. No. 2.9e-26;
                                                                                               Score 433; DB 8;
Pred. No. 2.9e-26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 433; DB 8;
Pred. No. 2.9e-26;
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Pred. No. 2.9e-26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22.7%; Score 433; DB 8; 33.8%; Pred. No. 2.9e-26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 433; DB 8;
Pred. No. 2:9e-26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABU12047 standard; protein; 422 AA.
Human NOV5a CG59883-01 protein SEQ ID 14.
WO200281625-A2.
 ADH43713 standard; protein; 352 AA.
Human PRO polypeptide #140.
US2003224984-A1.
                                                                                                                                                                                                                                                                                             ADI35682 standard; protein; 352 AA.
Human PRO polypeptide #108.
US2003096960-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                              ADH69776 standard; protein; 352 AA.
Human PRO polypeptide #108.
US2004019183-A1.
29-JAN-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADK83058 standard; protein; 352 AA.
Human PRO polypeptide #140.
US2004043927-A1.
04-MAR-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADK66692 standard; protein; 352 AA.
Human PRO polypeptide #108.
US2004044180-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW69698 standard; protein; 352 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US200401...
29-JAN-2004.
(GETH ) GENENTECH INC.
22.7%; SC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US200404...
04-MAR-2004.
(GETH ) GENENTECH INC.
22.7%; St.
                                             USZUCZ-

04-DEC-2003.

(GETH ) GENENTECH INC.

22.7%;
                                                                                                                                                                                            USZUCO-C.
22-MAY-2003.
(GETH ) GENENTECH INC.
22.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-MAR-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-OCT-2002.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 625
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                              Query Match
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Length 431;

Length 426;

Length 426;

Length 431;

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US2003092068-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 649
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 650
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LG.5%; Score 352.5; DB 8; Length 428;

LG.5%; Score 352.5; DB 8; Length 428;

LU ADP29312 standard; protein; 330 AA.

DE Human secreted protein SEQ ID #99.

PM WO20404035732-A2.

PD 29-APR-2004.

PA (FIVB-) FIVE PRIME THERAPEUTICS INC.

Query Match

Best Local Similarity 27.3%; Pred. No. 1.2e-19;

RESULT 638

ID AAW82731 standard; protein; 397 AA.

DE Adenovirus pACSG2SCAR.sig chime***

PM WO9854446-A1.

PM WO9854446-A1.

PA (FERTAL FOR THE PROTEIN FOR THE P
                                                                                                                                                                                                                                                                                                                                                                                                 AD657140 standard; protein; 428 AA.
Mouse B7-H5 protein SEQ ID NO:10.
W02004022594-A2.
18-MAT-2004.
(CYTO-) CYTOS BIOTECHNOLOGY AG.
ery Match
st Local Similarity 29.4%; Pred. No. 1.2e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 18.4%; Score 350.5; DB 2; Length 397; Local Similarity 28.2%; Pred. No. 1.5e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18.0%; Score 342.5; DB 2; Length 264; 33.9%; Pred. No. 3.9e-19;
                                                                                                                                                                                                  PD 31-JAN-2002.

PA (GETH) GENENTECH INC.

Query Match

Best Local Similarity 31.5%; Pred. No. 3e-20;

RESULT 636
                                   18.9%; Score 360; DB 8; Length 431; 30.2%; Pred. No. 2.9e-20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17.9%; Score 341; DB 2; Length 277; 34.6%; Pred. No. 5.5e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 341; DB 6; Length 412;
Pred. No. 9.4e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DACESTON OF THE PHARM INC. (MILL-) MILLENNIUM PHARM INC. (MILL-) MILLENNIUM PHARM INC. 18.3%; DB 4; Length 65; ery Match 18.3%; Score 350; DB 4; Length 65; ery Match 100.0%; Pred. No. 1.4e-20;
                                                                                                                               AAU83699 standard; protein; 290 AA. Human PRO protein, Seq ID No 216. WO200208288-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW82729 standard; protein; 264 AA.
Adenovirus pACTSG2-SCAR protein.
WO9854346-AI.
03-DEC-1998.
(GENV-) GENVEC INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW82730 standard; protein; 277 AA. Adenovirus SCAR.RGD protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADASO170 standard; protein; 412 AA.
Human CAR/SCF fusion protein.
US2003092068-A1.
US-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADA50171 standard; protein; 412 AA.
Human CAR/mouse SCF fusion protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB48124 standard; peptide; 65 AA.
Human A236 immunoglobulin domain.
WO200069885-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17.9%;
34.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
RESULF 6482730 standard; p:
DE Adenovirus SCAR.RGD pN WOS985436-A1.
PD 03-DEC-1998.
PA (GENV-) GENVEC INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ITOH/) ITOH A.
(HANA/) HANAZONO Y.
(OKAD/) OKADA T.
(OZAW/) OZAWA K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
RESULT 640
                                                        Best Local Similarity RESULT 635
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-DEC-1998.
(GENV-) GENVEC INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
RESULT 642
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
(UYTY ) UNIV TOKYO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ouery Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                   Query Match
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PD 25-FEB-2003.

PA (RAIN-) RAINBOW THERAPEUTIC CO.

Query Match 17.8%; Score 340.5; DB 6; Length 466;

Best Local Similarity 27.0%; Pred. No. 1.2e-18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MILL-) MILLENNIUM PHARM INC.

(MILL-) MILLENNIUM PHARM INC.

Lery Match

16.8%; Score 321; DB 4; Length 365;

Lery Match

28.4%; Pred. No. 3.3e-17;
                                                                                                                                                                                                                                                                                                                                17.9%; Score 341; DB 6; Length 493; 34.6%; Pred. No. 1.2e-18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17.6%; Score 335; DB 6; Length 474; 35.4%; Pred. No. 3.5e-18;
                                                                                                     Length 412;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17.6%; Score 335; DB 6; Length 393; 35.4%; Pred. No. 2.7e-18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 393;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADASO159 standard; protein; 474 AA.
Human CAR/mouse anti-CD34 antibody mature fusion protein.
US2003092068-A1.
                                                                                                                                                         ADA50172 standard; protein; 493 AA.
Human CAR/mouse anti-CD34 antibody fusion protein.
US2003092068-A1.
                                                                                                       17.9%; Score 341; DB 6; 34.6%; Pred. No. 9.4e-19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17.6%; Score 335; DB 6; 35.4%; Pred. No. 2.7e-18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADAS0158 standard; protein; 393 AA.
Human CAR/mouse SCF mature fusion protein.
US2003092068-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB65841 standard; protein; 365 AA.
Murine mature INTERCEPT 258 SEQ ID NO: 40.
WQ200078808-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADASO157 standard; protein; 393 AA. Human CAR/SCF mature fusion protein. US2003092068-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB65840 standard; protein; 394 AA.
Murine INTERCEPT 258 SBQ ID NO: 39-
WOZO0078808-A1.
28-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                            ABU62399 standard; protein; 466 AA
Chimeric CAR/Hg/Pro-A gene product
15-MAY-2003.
(ITOH/) ITOH A.
(HANA/) HANAZONO Y.
(OKAD/) OKADA T.
(OZAW/) OZAWA K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-MAY-2003.
(ITOH/) ITOH A.
(HANA) HANAZONO Y.
(OKAD/) OKADA T.
(OZAW/) OZAWA K.
                                                                                                                                                                                                                              15-MAY-2003.
(ITOH/) ITOH A.
(HANA/) HANAZONO Y.
(CKAD/) OKADA T.
(OZAW/) OZAWA K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      15-MAY-2003.
(ITOH/) ITOH A.
(HANA) HANAZONO Y.
(OKAD/) OKADA T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ITOH/) ITOH A.
(HANA/) HANAZONO Y.
(OKAD/) OKADA T.
(OZAW/) OZAWA K.
                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
RESULT 645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                     Query Match
Best Local Similarity
RESULT 644
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Best Local Similarity
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Length 327;

Length 327;

Length 327;

Length 327;

DB 6;

Length 327;

Length 327;

Length 327;

Length 327;

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Human membrane bound receptor/protein PR07154 amino acid sequence. US2003065147-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 309.5; DB 6; Length 327;
Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABOUJ1822 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154
US203045687-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel human secreted and transmembrane protein PRO7154 US2001088803-A1.
                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 30.2%; Pred. No. 2.4e-16;
RESULT 662
                                                                                                                                 GP-WOV-2001.

(HUMA-) HUMAN GENOME SCI INC.

(HUMA-) HUMAN GENOME SCI INC.

16.2%; Score 309.5; DB 5;

Lery Match

16.2%; Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 309.5; DB 6;
Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 309.5; DB 6;
Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16.2%; Score 309.5; DB 6; 30.2%; Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABO34368 standard; protein; 327 AA.
Human secreted/transmembrane polypeptide PRO 7154
US2003044934-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16.2%; Score 309.5; DB 6; 30.2%; Pred. No. 2.4e-16;
                     16.2%; Score 309.5; DB 4 30.2%; Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 309.5; DB 6
Pred. No. 2.4e-16;
                                                                          ABB90354 standard; protein; 327 AA. Human polypeptide SEQ ID NO 2730. WO200190304-AZ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein; 327 AA.
                                                                                                                                                                                                                                         AAU83709 standard; protein; 327 AA.
Human PRO protein, Seg ID No 236.
WO200208288-A2.
                                                                                                                                                                                                                                                                                                                                                                                                           ABU80856 standard; protein; 327 AA.
Human PRO polypeptide #118.
US2003036635-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein; 327 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein; 327 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 30.2%; P
RESULT 666
ID ABJ72473 standard; protein; 32°
DE Human PRO7154 protein.
PN US2003027988-A1.
PD 06-FEB-2003.
PA (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PD 08-MAY-2003.

PA (GETH ) GENENTECH INC.

QUENY MATCh 16.2%;

Best Local Similarity 30.2%;

RESULT 665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PD 06-MAR-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 30.2%;
RESULT 664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-FEB-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                         31-JAN-2002.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-MAR-2003.
(GETH ) GENENTECH INC.
    (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABJ72345 standard; pro
Human PRO7154 protein.
US2003050448-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
RESULT 663
                                                                                                                                                                                  Query Match
Best Local Similarity
RESULT 661
PA (GETH ) GENENTECH I
Query Match
Best Local Similarity
RESULT 660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABJ72175 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ouery Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADF83097 standard; protein; 326 AA.
Human corticol thymocyte receptor CTXL, overexpressed in cancer.
WO2003100000-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16.5%; Score 314.5; DB 4; Length 325; 29.4%; Pred. No. 9.3e-17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 327;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16.5%; Score 315.5; DB 8; Length 326; 30.0%; Pred. No. 7.8e-17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY87251 standard; protein; 327 AA.
Human signal peptide containing protein HSPP-28 SEQ ID NO:28.
WO200000610-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 309.5; DB 3; Length 327, Pred. No. 2.4e-16;
                     Length 394;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16.6%; Score 317; DB 4; Length 394; 28.2%; Pred. No. 7.6e-17;
                                                                                                                                                                                  Length 394;
                                                                                                                                                                                                                                                                                                                                                  16.8%; Score 320; DB 4; Length 394; 28.4%; Pred. No. 4.4e-17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 394
                                                                            AAB65908 standard; protein; 394 AA.
Murine secreted protein related protein SEQ ID NO: 142.
WO200078808-A1.
                                                                                                                                                                                                                                         AAB65910 standard; protein; 394 AA.
Murine secreted protein related protein SEQ ID NO: 146
W0200078808-A1.
28-DEC-2000.
(MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                           AAB65911 standard; protein; 394 AA.
Murine secreted protein related protein SEQ ID NO: 148.
WO200078808-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB65909 standard; protein; 394 AA.
Murine secreted protein related protein SEQ ID NO: 144
WO200078808-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 309.5; DB 3;
Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WG4000----
28-DE-2000.
(MILL-) MILLENNIUM PHARM INC.
(MILL-) MILLENNIUM PHARM INC.
ery Match
16.8%; Score 320; DB 4;
ery Match
28.4%; Pred. No. 4.4e-17;
                     16.8%; Score 321; DB 4; 28.4%; Pred. No. 3.6e-17;
                                                                                                                                 28-DEC-2000.
(MILL-) MILLENNIUM PHARM INC.
(MILL-) MILLENNIUM PHARM INC.
16.8%; Score 320; DB 4;
(ery Natch 13 4; Pred. No. 4.4e-17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB61480 standard; protein; 325 AA.
Human CTH protein.
WO200100672-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAY94857 standard; protein; 327 AA.
Human protein clone HP10568.
WO200005367-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY97585 standard; protein; 327 AA. Human secreted protein PRO7154. WO200075317-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-DEC-2000.
(MILL-) MILLENNIUM PHARM INC.
(MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-JAN-2001
(MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-FEB-2000.
(SAGA ) SAGAMI CHEM RES CENT.
(PROT-) PROTEGENE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (INCY-) INCYTE PHARM INC
                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
RESULT 654
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-DEC-2003.
(TULA-) TULARIK INC.
              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                             Query Match
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Best Local Similarity 30.2%; Pred. No. 2.4e-16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1D ADB83972 standard; protein; 327 AA.

DE Novel human secreted and transmembrane protein PRO7154.

DE Novel human secreted and transmembrane protein PRO7154.

PN US2003069397-A1.

PD 10-APR-2003.

PA (GETH ) GENENTECH INC.

16.2%; Score 309.5; DB 7; Length 327;

Query Match
   Score 309.5; DB 7; Length 327; Pred. No. 2.4e-16;
                                                                                                                                                                                                                                             UB-TMAI-2000.
(GETH ) GENENTECH INC.
iry Match 16.2%; Score 309.5; DB 7; Length 327;
ir foral Similarity 30.2%; Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 309.5; DB 7; Length 327; Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                      Length 327;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 309.5; DB 7; Length 327; Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 327;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 309.5; DB 7; Length 327;
Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 309.5; DB 7; Length 327;
Pred. No. 2.4e-16;
                                                                                                                                      Length 327;
                                                                                                                                                                                    ADB80832 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154.
US2003088068-A1.
08-MAY-2003.
                                                                                                                                                                                                                                                                                                             ADB73373 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154
US2003096968-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel human secreted and transmembrane protein PRO7154. Usus 201929886-Al.
                                                                    Novel human secreted and transmembrane protein PRO7154 US2003073814-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADB/88455 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154
US2003092889-A1.
                                                                                                                                    Score 309.5; DB 7;
Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                      Score 309.5; DB 7;
Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 309.5; DB 7;
Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human PRO polypeptide #118.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human PRO polypeptide #118.
US2003088067-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB85103 standard; protein; 327 AA. Human PRO polypeptide #118. US2003073817-A1. 17-APR-2003. (GETH ) GENENTECH INC.
                                                   protein; 327 AA
                                                                                    US20030202
22-MAY-2003.
(GETH ) GENENTECH INC.
16.2%; F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          USZUOSUC
08-MAY-2003.
(GETH ) GENENTECH INC.
MATCh 13-rity 30.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Match 16.2%; Local Similarity 30.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GETH ) GENENTECH INC.
Query Match
RESULT 669
ID ADB83726 standard; pz
DE Novel human secreted
Pp USCO3027384-A1.
PD 17-ARR-2003.
PA (GETH ) GENENTECH INC
                                                                                                                                              Best Local Similarity RESULT 670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity RESULT 676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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Best Local Similarity
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Score 309.5; DB 7; Length 327; Pred. No. 2.4e-16;
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                                                                                                                                                                                                                                                                                   Score 309.5; DB 7; Length 327; Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 7; Length 327;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 309.5; DB 7; Length 327;
Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 309.5; DB 7; Length 327;
Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 327;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 309.5; DB 7; Length 327;
Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16.2%; Score 309.5; DB 7; Length 327; 30.2%; Pred. No. 2.4e-16;
                ADB73127 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154.
US2003092887-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AUC47563 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154.
US200308072-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADC47308 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154.
US2003105288-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADC49986 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154
US2003088064-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC49702 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154 US2003088071-A1.
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Novel human secreted and transmembrane protein PRO7154
US2003096972-A1.
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Novel human secreted and transmembrane protein PRO7154
US2003088070-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                             16.2%; Score 309.5; DB 7 30.2%; Pred. No. 2.4e-16;
                                                                                                                                                                                                                                                                                                                                 Human PRO polypeptide #118.
US2003096969-A1.
                                                                                                                                                                          ADC36965 standard; protein; 327 AA.
Human PRO polypeptide #118.
US2003088065-A1.
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RESULT 680
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                                                                                                                 Query Match 16.2%;
Best Local Similarity 30.2%;
RESULT 679
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Best Local Similarity 30.2%;
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Best Local Similarity 30.2%;
RESULT 685
                                                                             15-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
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RESULT 678
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ADE05954 standard; protein; 327 AA.
Human PRO polypeptide #118.
US2003100728-A1.
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Pred. No. 2.4e-16;
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Pred. No. 2.4e-16;
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Pred. No. 2.4e-16;
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Pred. No. 2.4e-16;
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                 ADD06418 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154.
US2003073816-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADD51146 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154.
US2003105290-A1.
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Novel human secreted and transmembrane protein PRO7154.
US2003105289-A1.
OS-JUN-2003.
(GETH.) GENENTECH INC.
                                                                                                                                                                 ADC77937 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154
US2003088066-A1.
                                                                                                                                                                                                                                                                                                                ADD50900 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154
US2003105291-A1.
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Novel human secreted and transmembrane protein PRO7154.
US2003100735-A1.
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Novel human secreted and transmembrane protein PRO7154
US203092888-A1.
                                                                                                              DB 7;
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Pred. No. 2.4e-16;
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Pred. No. 2.4e-16;
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Pred. No. 2.4e-16;
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17-APR-2003.
(GETH ) GENENTECH INC.
""rch 16.2%;
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RESULT 694
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Human PRO polypeptide #118.
US2003096970-A1.
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Best Local Similarity 30.2%;
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Local Similarity 30.2%;
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US2003096971-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                        Best Local Similarity RESULT 688
                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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Best Local Similarity
RESULT 696
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RESULT 687
ID ADD064
DE Novel
PN US2003
PD 17-APR
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Best Local Similarity 30.2%; Score 309.5; DB 8; Length 327; RESULT 703
DE Human pro ...
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                                                 Length 327;
                                                                                                                                                                                                           Length 327;
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Pred. No. 2.4e-16;
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Novel human secreted and transmembrane protein PRO7154
US2003100717-A1.
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Novel human secreted and transmembrane protein PRO7154.
US2003100738-A1.
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Novel human secreted and transmembrane protein PRO7154
US2003100722-A1.
29-MAY-2003.
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Novel human secreted and transmembrane protein PRO7154
US2003100734-A1.
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Novel human secreted and transmembrane protein PRO7154
US2003096362-A1.
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                                                 8;
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Pred. No. 2.4e-16;
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Pred. No. 2.4e-16;
                                                                                                                                                                                                           Score 309.5; DB 8;
Pred. No. 2.4e-16;
                                                 Score 309.5; DB 8
Pred. No. 2.4e-16;
                                                                                                   DE Human PRO polypeptide #118.

PRO US2003100712-A1.

PD 29-MAY-2003.

PA (GETH ) GENENTECH INC.

Query Match

Best Local Similarity 30.2%; Pred. No

RESULT 698

ID ADD75929 standard; protein; 327 AA.

DE Novel human secreted and transmembrance

PRO US203100717-A1.

PRO US2003100717-A1.

PRO US2003100717-A1.

PRO GETH ) GENENTECH INC.
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SULT 704
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PD 29-MAY-2003.
PA (GETH ) GENENTECH INC.
QUERY MATCh
Best Local Similarity 30.2%;
RESULT 697
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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US2003100726-A1.
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RESULT 723
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Pred. No. 2.4e-16;
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Pred. No. 2.4e-16;
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Pred. No. 2.4e-16;
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                                                                                                                                                                                                                                                                          ADE21356 standard; protein; 327 AA.

Novel human secreted and transmembrane protein PRO7154
US2003100736-A1.
29-MAY-2003.
(GETH ) GENENTECH INC.
16.2%; Score 309.5; DB 8; Lengt
Bt Local Similarity 30.2%; Pred. No. 2.4e-16;
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Novel human secreted and transmembrane protein PRO7154.
US2003100733-A1.
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Novel human secreted and transmembrane protein PRO7154.
US2001100718-A1.
29-MAY-2003.
(GETH) GENENTECH INC.
16.2%; Score 309.5; DB 8; Lengt Local Similarity 30.2%; Pred. No. 2.4e-16;
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Novel human secreted and transmembrane protein PRO7154
US2003100732-A1.
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NOVEL human secreted and transmembrane protein PRO7154
US2003100721-A1.
29-MAY-2003.
(GETH ) GENENTECH INC.
                           Novel human secreted and transmembrane protein PRO7154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 309.5; DB 8;
Pred. No. 2.4e-16;
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Pred. No. 2.4e-16;
                                                                                                                                                                               Score 309.5; DB 8;
Pred. No. 2.4e-16;
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Human PRO polypeptide #118.
US2003100064-A1.
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Human PRO polypeptide #118.
US2003100709-A1.
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Human PRO polypeptide #118.
Novel human ...
US2003100737-A1.
29-MAY-2003.
(GETH ) GENENTECH INC.
***rch 16.2%; Sc
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(GETH ) GENENTECH INC.
Watch 130.2%; Sr
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29-MAY-2003.
(GETH ) GENENTECH INC. .
MATCh ... 1 arity 30.2%;
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29-MAY-2003.
(GETH ) GENENTECH INC.
Match ''rrity 30.2%;
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RESULT 711
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human PRO polypeptide #118
US2003100708-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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J.04-A1.
2003.
LOG3.
LY MATCH
BEST LOCAL SIMILARITY
RESULT 710
ID ADD74199 stand<sup>2</sup>
ID A
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Best Local Similarity
RESULT 707
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Best Local Similarity
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Best Local Similarity
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Score 309.5; DB 8; Length 327; Pred. No. 2.4e-16;
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                                                                                                                                                                                                                                                      ADD76973 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154
US2003100715-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                    ADD86741 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154
US2003100719-A1.
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Novel human secreted and transmembrane protein PRO7154
US2003100731-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AD077717 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154
US2003100729-A1.
29-MAY-2003.
(GETH ) GENENTECH INC.
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Novel human secreted and transmembrane protein PRO7154
US2003100725-A1.
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Novel human secreted and transmembrane protein PRO7154
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                                                                                             ADD75429 standard; protein; 327 AA. Human PRO polypeptide #118.
US2003100714-Al. 29-MAY-2003.
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US2003100710-A1.
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US2003100713-A1.
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29-MAY-2003.
(GETH ) GENENTECH INC.
16.2%;
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(GETH) GENENTECH INC.

Query Match
Best Local Similarity 30.2%;
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                    GENENTECH INC.
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(GETH ) GENENTECH INC
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Best Local Similarity
RESULT 719
                                       Query Match
Best Local Similarity
RESULT 715
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29-MAY-2003
(GETH ) GEN
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Length 327;

DB 8;

Length 327;

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DB 8;

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Score 309.5; DB 8; Length 327; Pred. No. 2.4e-16;
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Human colon cancer antigen protein sequence SEQ ID NO:847.
WO200055351-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AD129957 standard, protein, 327 AA.
Novel human secreted and transmembrane protein PRO7154.
US2003096961-A1.
                                                                                                                                                                                  Annay142 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154
US2003096655-A1.
                                                                                                                                                                                                                                                                                                                   AUG34232 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154
US2004006206-A1.
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Novel human secreted and transmembrane protein PRO7154
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Pred. No. 2.4e-16;
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Pred. No. 2.4e-16;
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Pred. No. 2.4e-16;
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29-TAN 20-TAN A.
                                                            ADG06798 standard; protein; 327 AA. Human PRO polypeptide #118.
US2003095966-A1.
2-MAY-2003.
(GETH ) GENENTECH INC.
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(HUMA-) HUMAN GENOME SCI INC.
               16.2%;
30.2%;
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(GETH) GENENTECH INC.
Query Match
16.2%;
Best Local Similarity 30.2%;
SULT 737
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(GETH) GENENTECH INC.
Query Match 16.2%;
Best Local Similarity 30.2%;
SULT 738
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US2004044180-A1.
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(GETH ) GENENTECH INC.
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GENENTECH INC
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                                                                                                                                                            Best Local Similarity RESULT 734
                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 735
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             Query Match
Best Local Similarity
RESULT 733
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Best Local Similarity
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ry Match

16.2%; Score 309.5; DB 8; Length 327;

r nocal Similarity 30.2%; Pred. No. 2.4e-16;
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Pred. No. 2.4e-16;
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Novel human secreted and transmembrane protein PRO7154.
US2003096964-A1.
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Novel human secreted and transmembrane protein PRO7154
US2003096959-A1.
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Novel human secreted and transmembrane protein PRO7154
US2003096967-A1.
                                                                               ADD77219 standard; protein; 327 AA.
Novel human secreted and transmembrane protein PRO7154
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(GETH) GENENTECH INC.

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cery Match

16.2%; Score 309.5; DB 8;
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Pred. No. 2.4e-16;
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Pred. No. 2.4e-16;
                                Score 309.5; DB 8;
Pred. No. 2.4e-16;
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19200310724-Al.
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Human PRO polypeptide #118.
US2003100723-A1.
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22-MAY-2003.
(GETH ) GENENTECH INC.
43-FCh 16.2%; S'
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29-MAY-2003.

A (GETH) GENENTECH INC.

Query Match

Best Local Similarity 30.2%;
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Human PRO polypeptide #118.
US2003096962-A1.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                    29-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
Query Match 1
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Best Local Similarity
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                                                                                                                                                                                    Local Similarity
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Best Local Similarity
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                                                                                                                    US2003100716-A1.
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                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                      Query Match
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RESULT 750
ID ADP54587 standard; protein; 319 AA.
DE Human PRO protein sequence SEQ ID NO:563.
PN WO2004039956-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ALPH-) ALPHAGENE INC.
Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RUBEN S M.
FLORENCE K A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (BREW) BREWER L A.
(SOPP) SOPPET D R.
(LAFL/) LAFLEUR D W.
(ENDR/) ENDRESS G A.
(EBNE/) EBNER R.
(BIRS/) BIRSE C E.
                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity RESULT 754
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NI J.
ROSEN C A.
CARTER K C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOORE P A.
OLSEN H S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        YOUNG P E.
                                                                                                                                                                                                                       07-MAR-2002.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NIJJ/)
ROSE/)
CART/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        YOUN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MOOR/)
(OLSE/)
                                                                                                                                                 RESULT 751
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Loc
RESULT 756
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADN'99847 standard; protein; 319 AA.
Cancer/angiogenesis/fibrosis-related polypeptide, SEQ ID NO:C217.
WO2003042661-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADA10947 standard; protein; 319 AA.
Human cDNA differentially expressed in colon cancer #43 product.
US2002160382-A1.
                                                                                                                                                                                                                                                   06-MAR-1997.

1 (LUDW-) LUDWIG INST CANCER RES.

Query Match

16.2%; Score 308.5; DB 2; Length 319;

Ouery Match

16.2%; Pred. No. 2.8e-16;
                                                                                                                                                                                                                                                                                                                                                                                                 03-JUN 1999.
(GETH ) GENENTECH INC.
ery Match 16.2%; Score 308.5; DB 2; Length 319;
ery Match 31.2%; Pred. No. 2.8e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IS-APR-2004.

(GETH ) GENENTECH INC.

Query Match

16.2%; Score 308.5; DB 8; Length 319;

Query Match

16.2%; Pred. No. 2.8e-16;
                                                                                                      OS-APR-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match
ery Match
16.2%; Score 309.5; DB 4; Length 365;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-DEC-2000.
(MILL-) MILLENNIUM PHARM INC.
ery Match
st Local Similarity 31.2%; Pred. No. 2.8e-16;
 16.2%; Score 309.5; DB 3; Length 365; 30.2%; Pred. No. 2.8e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 308.5; DB 6; Length 319;
Pred. No. 2.8e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-MAY-2003.
(ROSB.) EOS BIOTECHNOLOGY INC.
16.2%; Score 308.5; DB 7; Length 319;
rt Local Similarity 31.2%; Pred. No. 2.8e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 319
                                                        protein; 365 AA.
antigen protein SEQ ID NO:6377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16.2%; Score 308.5; DB 7; 31.2%; Pred. No. 2.8e-16;
                                                                                                                                                                                                                                                                                                                                                  AAY23323 standard; protein; 319 AA.
Amino acid sequence of the A33 antigen.
WO9927098-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB65863 standard; protein; 319 AA.
Human A33 protein SEQ ID NO: 67.
WO200078808-A1.
                                                                                                                                                                                                    AAW14146 standard; protein; 319 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein; 319 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADH62533 standard; protein; 319 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16.2%;
31.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human A33 antigenic protein
US2003171568-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A. (135289 standard; protein
Human A33 antigen protein.
WO2004031105-A2.
               Best Local Similarity
RESULT 742
ID AAG75613 standard; pr
DE Human colon cancer ar
PN WO200122920-A2.
PD 05-APR-2001.
PA (HUMA-) HUMAN GENOME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (FONG)) FONG S.
(GURN) GURNEY A L.
(IMPL) NAPIER M A.
(TUMA/) TUMAS D.
(WODD) WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ASHKENAZI A.
                                                                                                                                                          Best Local Similarity RESULT 743
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                         Human A33 antigen.
WO9708189-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (LASE/) LASEK A W. (JONE/) JONES D A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-SEP-2003.
(ASHK/) ASHK
(FONG/) FONG
(GODD/) GODD/
(GURN/) GURN/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Loca
RESULT 748
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DB 8; Length 319;
                                                                                                                                                                                          16.2%; Score 308.5; DB 5; Length 336; 31.2%; Pred. No. 3e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 307.5; DB 2; Length 390; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 389;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 389;
                                                                                                                                                                                                                                                                                                                                           Length 316;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 325;
                                                                                                                                                                                                                                                                                                                                                                                            AAY95024 standard; protein; 325 AA.
Human clone vc51_1 deletional variant ORF1, SEQ ID NO:124.
WO200011015-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAY76303 standard, protein, 389 AA.
Fragment of human secreted protein encoded by gene
WO9958660-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16.1%; Score 307.5; DB 3; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 307.5; DB 7;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                    Luery Match 16.1%; Score 308; DB 8; Best Local Similarity 31.1%; Pred. No. 3e-16; RESULT 75.00 Standard; protein; 325 AA DE Human clone vc51_1 deletion.

PN W0200011015-A1.

PA 02-MAR-200.
                                    Score 308.5; DB 8
Pred. No. 2.8e-16;
                                                                                                                                                                                                                                                                                                                                         DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16.1%; Score 308; DB 3; 29.8%; Pred. No. 3.1e-16;
                                                                                 Human polypeptide SEQ ID NO 318. WO200218424-A2.
                                                                                                                                                                                                                                            ADK40854 standard, protein; 316 AA.
Human A33 molecule.
US6699688-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADE11956 standard; protein; 389 AA.
Human secreted polypeptide #210.
US2003100051-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAY05286 standard; protein; 390 AA.
EGF-like homologue PRO246.
WO9914327-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-NOV-1999.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PD 25-MAR.1999.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 29.2%;
) 13-MAY-2004.

(GETH ) GENENTECH INC.

Query Match 16.2%;

BEST Local Similarity 31.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16.1%;
29.2%;
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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 772
                                                                             Score 307.5; DB 2; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                        Score 307.5; DB 2; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 16.1%; Score 307.5; DB 3; Length 390; Best Local Similarity 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 307.5; DB 4; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16.1%; Score 307.5; DB 4; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16.1%; Score 307.5; DB 4; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                   Score 307.5; DB 3; Length 390; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 307.5; DB 4; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 307.5; DB 4; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AABS3082 standard; protein; 390 AA.
Human angiogenesis-associated protein PRO246, SEQ ID NO:96.
WO200053753-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAB99818 standard; protein; 390 AA.
Human shear stress-response protein SEQ ID NO: 144
WO200125427-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAB31207 standard; protein; 390 AA.
Amino acid sequence of human polypeptide PRO246.
WO200077037-A2.
21-DEC-2000.
                                                                                                                                                                                                                                                                                              AAY94999 standard; protein; 390 AA.
Human secreted protein vc51_1, SEQ ID NO:38.
WO200011015-A1.
Amino acid sequence of protein PRO246.
                                                                                                                                 AAY27096 standard; protein; 390 AA. Human viral receptor protein (ACVRP) US5942606-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY88574 standard; protein; 390 AA.
Human PRO246 amino acid seguence.
WO200015666-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAU12340 standard; protein; 390 AA.
Human PRO246 polypeptide sequence.
WO200140466-AZ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein; 390 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAB68599 standard; protein; 390 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                W020010-2-18-JAN-2001.
(GETH ) GENENTECH INC.
"artch "artch 29.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-APR-2001.
(KYOW ) KYOWA HAKKO KOGYO KK.
(NOJI/) NOJIMA H.
                                                                                                                                                                                                                                    16.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 29.2%;
RESULT 765
                                                                       Best Local Similarity 29.2%;
RESULT 758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 16.1%;
Beet Local Similarity 29.2%;
RESULT 766
                                                                                                                                                                                                                                                                                                                                                           02-MAR-2000.

(ALPH-) ALPHAGENE INC.

Query Match

Best Local Similarity 29.2%;
                                                                                                                                                                                              24-AUG-1999.
(INCY-) INCYTE PHARM INC.
                                   25-MAR-1999.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-MAR-2000.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-JUN-2001.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-SEP-2000.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AABB0219 standard; pr
Human PRO246 protein.
WO200104311-Al.
                                                                                                                                                                                                                                      Query Match
Best Local Similarity
RESULT 759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity RESULT 762
                  WO9914328-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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DB 4; Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4; Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 6; Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6; Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ouery Match 16.1%; Score 307.5; DB 6; Length 390; Best Local Similarity 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16.1%; Score 307.5; DB 6; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABO17784 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003032156-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABO01781 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AB021178 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003040014-A1.
                                                                                                                                                                                                                                                                          AAB88358 standard; protein; 390 AA.
Human membrane or secretory protein clone PSEC0086.
EP1067182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16.1%; Score 307.5; DB 6; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 29.2%; Pred. No. 4.4e-16;
RESULT 770
                                                                                                                                                                    Score 307.5; DB 4
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16.1%; Score 307.5; DB 4 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16.1%; Score 307.5; DB 6 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 307.5; DB 6
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16.1%; Score 307.5; DB 6 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABU71898 standard; protein; 390 AA.
Human secreted/transmembrane protein PRO246.
US2003003530-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human PRO polypeptide #169.
US2003004311-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABU71452 standard; protein; 390 AA.
Human PRO polypeptide #8.
US2002192659-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein; 390 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA56949 standard; protein; 390 AA
PRO246.
WO200105836-A1.
J 25-JAN-2001.
A (GETH) GENENTECH INC.
A "** CAT "** C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PD 10-0CT-2002.
PA (GETH ) GENENTECH INC.
QUERY Match 16.1%;
Best Local Similarity 29.2%;
RESULT 769
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human secreted protein #232.
WO2002102994-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human PRO polypeptide #8.
US2002146709-A1.
                                                                                                                                                                                                                                                                                                                                                                                           10-JAN-2001.
(HELI-) HELIX RES INST.
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(GETH.) GENENTECH INC.
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(GETH ) GENENTECH INC
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                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 767
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
RESULT 768
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABU71597 standard;
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16.1%;
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FESULT 788
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RESULT 787
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Best Local Similarity 29.2%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                        28-NOV-2002.
(GETH ) GENENTECH INC.
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                                                      Best Local Similarity
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Best Local Similarity
RESULT 786
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US2003032057-A1.
           31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-FEB-2003
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                                          Query Match
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                                                         Score 307.5; DB 6; Length 390; Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                     16.1%; Score 307.5; DB 6; Length 390; 29.2%; Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABU67296 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003032063-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    16.1%; Score 307.5; DB 6; 29.2%; Pred. No. 4.4e-16;
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Human secreted/transmembrane polypeptide PRO246.
US2003044839-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABUS9819 standard; protein; 390 AA.
Novel secreted and transmembrane protein PRO246.
US2003017563-A1.
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Human secreted/transmembrane protein (PRO) #169.
US2003036179-A1.
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                                                                                                                                                                                                                                    ABU54354 standard; protein; 390 AA.
Human secreted/transmembrane protein PRO246.
US2002132240-A1.
19-SEP-2002.
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Human secreted/transmembrane protein, #9
US2002160374-A1.
                                                                                                  ABU66738 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003036180-AI.
CO-FEB-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           D ABR47754 standard, protein, 390 AA.
E Human secreted protein, SEQ ID 645.
N W2020295010-A2.
A (HUMA-) HUMAN GENOME SCI INC.
Query Match
                                                                                                                                                                                                                                                                                                                                                                    ADA40800 standard; protein; 390 AA
                                                                                                                                                                                                                                                                                                                                                                                     Human secreted protein.
w02002102993.A2.
27-DEC-2002.
(HUMA-) HUMAN GENOME SCI INC.
        US200215...
26-DEC-2002.
(GETH ) GENENTECH INC.
16.1%;
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PA (GETH ) GENENTECH INC.
Query March
Best Local Similarity 29.2%;
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Query Match
Best Local Similarity 29.2%;
RESULT 784
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J.H.)

LH.) GENENTECH J.

LY MATCH
BEST LOCAL SIMILARITY
RESULT 776
ID ABUG6738 standa
DE Human PRO
PN US20030-
PN US20030-
PD 20-
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Best Local Similarity
RESULT 779
ID ABU67296 standard; pl
DE Novel human secreted
PN USZO03032063-A1.
PD 13-FEB-2003.
PA (GETH ) GENENTECH INC
                                                                                                                                                                                                Best Local Similarity RESULT 777
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RESULT 782
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31-UC1-Zave.
(GETH ) GENENTECH INC.
ry Match
16.1%; Score 307.5; DB 6; Length 390;
r Toral Similarity 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 6; Length 390;
                                                                                                                                                                                                              Length 390;
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Pred. No. 4.4e-16;
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Novel human secreted and transmembrane protein PRO246.
30-JAN-2003.
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Human secreted/transmembrane, PRO, protein SEQ ID 338.
US2003032155-A1.
                                                                                                ABU72064 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2002177165-A1.
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Novel human secreted and transmembrane protein PRO246.
US2003017463-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human secreted / transmembrane polypeptide PRO246.
US2003036060-A1.
20-FEB-2003.
(GETH ) GENENTECH INC.
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Human secreted / transmembrane polypeptide PRO246
US2003027143-A1.
                                                                                                                                                                                                              Score 307.5; DB 6;
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Human secreted/transmembrane protein PRO246.
                                                                                                                                                                                                                                                                              ABUG7352 standard, protein, 390 AA. Human secreted protein PRO246. US2003023054-A1. 30-JAN-2003. (GETH ) GENENTECH INC.
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Length 390;
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Novel human secreted and transmembrane protein PRO246.
US2003082693-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADA87469 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003087345-A1.
                                                                                                                                                                                 Score 307.5; DB 6;
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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                                                                                                                                                                                                                              AB032763 standard; protein; 390 AA.
Human secreted/transmembrane protein PRO246.
US2003045693-A1.
                                                                                                  ADA18100 standard; protein; 390 AA. Human secreted/transmembrane protein, #9. US2003039911-A1. 27-FBB-2003.
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01-Mav.ac.
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01-Mav.anc.
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US2003087349-A1.
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Human PRO polypeptide #169.
US2003068795-A1.
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Human PRO polypeptide #169.
US2003068794-Al.
                                                                                                                                                                                16.1%;
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RESULT 807
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RESULT 809
Luery Match
Best Local Similarity
RESULT 803
ID ADA.8100 stand>
DE Human secr'
PN US2003^-
PD 27-
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
RESULT 805
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Best Local Similarity
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                                                                                                                                                                                                  Score 307.5; DB 6; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 307.5; DB 6; Length 390;
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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                                                                     Score 307.5; DB 6; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                               Length 390;
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Pred. No. 4.4e-16;
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Pred: No. 4.4e-16;
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Pred. No. 4.4e-16;
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Novel human secreted and transmembrane protein PRO246.
US2003068796-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADA86366 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003082711-A1.
                                                                                                                                                                                                                                                                                                                               DB 6;
                                                                                                                                                                                                                                                                                                                             Score 307.5; DB 6
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                             ADB29244 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9
US2003092002-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human PRO polypeptide #169.
US2003082704-A1.
                                                                                                               ADA76288 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003073212-A1.
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08-mav_ana
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADA61561 standard; protein; 390 AA
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10.APR-2003.
(GETH ) GENENTECH INC.
Match ''rrity 29.2%;
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01-MX-2003.
(GETH ) GENENTECH INC.
16.1%;
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01-MY-2003.
(GETH ) GENENTECH INC.
16.11;
                                                                     16.18;
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RESULT 797
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ry Match 16.1%;
t Local Similarity 29.2%;
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Best Local Similarity 29.2%;
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US2003073215-A1.
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US2003049816-A1.
13-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                          20-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                               15-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                   (GETH ) GENENTECH INC.
                                                                     Query Match
Best Local Similarity
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Best Local Similarity
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RESULT 799

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01-MAY-2003
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     RESULT 821
                                                                                                                                                                                                                                                                                                                                                                 Ol-rwi-zvos.
(GETH ) GENENTECH INC.
rry Match
rry Match 16.1%; Score 307.5; DB 6; Length 390;
rry Annal Similarity 29.2%; Pred. No. 4.4e-16;
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(GETH) GENENTECH INC.
ery Match 16.1%; Score 307.5; DB 6; Length 390;
-- '-- Similarity 29.2%; Pred. No. 4.4e-16;
                                                                                                                       Score 307.5; DB 6; Length 390; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                            Length 390;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Novel human secreted and transmembrane protein PRO246.
US2003082691-A1.
                                                                                                                                                                                                                                                                                                        ADA91763 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003082694-A1.
01-MAY-2003.
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PRO246. USZOJSOJSJIL-AI.
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Human secreted/transmembrane polypeptide PRO246.
US2003044844-A1.
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Pred. No. 4.4e-16;
   29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                      ADA16075 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9
US2003049621-A1.
                          Human PRO polypeptide #8.
US2003044793-A1.
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Human PRO polypeptide #169.
US2003087351-A1.
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19200307722-A1.
                                                                  US200507.
06-MAR-2003.
(GETH ) GENENTECH INC.
March 16.1%; Sr
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08-MAY-2003.
(GETH ) GENENTECH INC.
16.1%; S'
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13-MARZ 2003.
(GETH ) GENENTECH INC.
16.1%;
ARCh -- 13-rity 29.2%;
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(GETH ) GENENTECH INC.
MATCh -11-rity 29.24;
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ry Match
t Local Similarity 29.2%;
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Local Similarity 29.2%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
Best Local Similarity RESULT 812
                                                                                                                       Query Match
Best Local Similarity
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Best Local Similarity
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Best Local Similarity
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Score 307.5; DB 6; Length 390;
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
ABO43317 standard; protein; 390 AA.
Novel human serreted and transmembrane protein PRO246.
US2003044945-A1.
(G6-WAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AMASSZ62 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
USZ003082695-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADA84710 standard; protein; 390 AA.

Novel human secreted and transmembrane protein PRO246.
US2003082708-A1.
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Pred. No. 4.4e-16;
                                                                                          Score 307.5; DB 6
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                                                         ADA42220 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003054401-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                    ADB24697 standard; protein; 390 AA. Human PRO polypeptide SEQ ID NO 338 US2003077713-A1.
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US2003064367-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADA75184 standard, protein; 390 AA.
Human PRO polypeptide #169.
US2003073216-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADA82221 standard; protein; 390 AA.
                                                                                                                                                ADA74464 standard; protein; 390 AA
                                                                                                                                                                                                                                        16.1%;
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                                                                                    Best Local Similarity 29.2%;
RESULT 822
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RESULT 829
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Best Local Similarity 29.2%;
RESULT 830
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US2003082701-A1.
                                                                                                                                                               Human PRO polypeptide #169.
US2003068798-A1.
                                                                                                                                                                                                    10-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                 20-MAR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 823
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Best Local Similarity
RESULT 824
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RESULT 825
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Human secreted/transmembrane protein, #9.
           Human PRO polypeptide SEQ ID NO 338.
US2003077714-A1.
24-APR-2003.
24-APR-2003.
EY MAICH
ELOCAL Similarity 29.2%; Pred. N
                                                                                                                                                                                                                                                                                            ADA81046 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003082702-A1.
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Human PRO polypeptide #169.
US2003082760-A1.
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Human PRO polypeptide #169.
US2003077710-A1.
                                                                                                                                           ADA96474 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003082690-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                           ADA95922 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003082759-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADA77495 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003068797-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADA16499 standard; protein; 390 AA
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29.2%;
                                                                                                                                                                                                                                                                                                                                          PD 01-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match 16.1%;
Best Local Similarity 29.2%;
RESULT 842
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 29.2%;
RESULT 841
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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                                                                                                    Best Local Similarity RESULT 840
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(GETH ) GEN
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                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 307.5; DB 6; Length 390;
Pred. No. 4.4e-16;
                                                                                                      Score 307.5; DB 6; Length 390;
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB25257 standard, protein, 390 AA.
Human PRO polypeptide SEQ ID NO 338
US2003077715-A1.
Human PRO polypeptide #169.
17-App-20.
                                                                                                                                                            ADA80494 standard, protein, 390 AA.
Human PRO polypeptide #169.
US2003082761-Al.
                                                                                                                                                                                                                                                                                                                   protein; 390 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA46961 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003073210-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA93433 standard, protein, 390 AA.
Human PRO polypeptide #169.
US2003077721-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB26783 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003092147-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB31070 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003096386-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADB24145 standard; protein; 390 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADA60998 standard; protein; 390 AA
                                                                                                                                                                                                        USACOZ-
01-MAZ-2003.
(GETH ) GENENTECH INC.
MATCh 114;
                                                                 PD 17-APR-2003.
PA (GETH ) GENENTECH INC.
Query Match 16.1%;
Best Local Similarity 29.2%;
RESULT 831
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Match 16.1%;
Local Similarity 29.2%;
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(GETH) GENENTECH INC. .
PRY MATCH 16.1%;
It Local Similarity 29.2%;
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Best Local Similarity 29.2%;
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Local Similarity 29.2%;
                                                                                                                                                                                                                                                                                                             ADA75736 standard, protein,
Human PRO polypeptide #169.
US2003082703-A1.
                                                                                                                                                                                                                                                                                                                                                                       01-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
RESULT 832
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
US2003049817-A1.
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DB 6; Length 390;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 307.5; DB 7; Length 390; Pred. No. 4.4e-16;
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  6; Length 390;
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Novel human secreted and transmembrane protein PRO246.
US2003082765-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Andrewick standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003082109-A1.
                                                                                                                                                                                                                                                                                                                                                                      DB 6;
Score 307.5; DB 6 Pred. No. 4.4e-16;
                                                                                                                                                                                  Score 307.5; DB 6
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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27-MAR-2003.
(GETH ) GENENTECH INC.
                                  Query Match
                                                                                                                                                                                                                                                                                                                                                         RESULT 860
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                                                                                                                                                                                           Score 307.5; DB 7; Length 390;
Pred. No. 4.4e-16;
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                                                   Score 307.5; DB 7; Length 390;
Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 390;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 7; Length 390;
                                                                                                                                                                                                                                                                                                                                                                                      ADA88021 standard; protein; 390 AA.

Novel human secreted and transmembrane protein PRO246.
US2003082700-A1.
01-MAY-2003.
(GETH) GENENTECH INC.
16.1%; Score 307.5; DB 7; Lengt Local Similarity 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel human secreted and transmembrane protein PRO246 US2003054516-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 307.5; DB 7;
Pred. No. 4.4e-16;
                                                                                               ADA12928 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003049622-A1.
                                                                                                                                                                                                                                         ADA41796 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
01-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADA17143 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003017498-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA42646 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9
US2003054351-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADB28439 standard, protein, 390 AA.
Human PRO polypeptide #169.
US2003082699-A1.
01-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADB28991 standard, protein; 390 AA.
Human PRO polypeptide #169.
US2003082706-A1.
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Human PRO polypeptide #169.
US2003059909-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADA46409 standard; protein; 390 AA
                                                                           "...ard; protein,
"...Ja04962-A1.

13-MAR-2003.

PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 29.2%; Pre
RESULT 850
ID ADA41796 standard; protein
PD Human secreted/tr-
PN US2003082547
PA /...
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01-MAY-2003.
(GETH) GENENTECH INC.
16.1%; SC
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20-MAR-2003.
(GETH ) GENENTECH INC.
Match 15.1%;
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23-JUN-2003.
(GETH ) GENENTECH INC.
MATCh ''nrity 29.2%;
                                                   16.18; 29.28;
                                                                                                                                                                                                                                                                                                                                     16.1%;
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PA (GETH ) GENENTECH INC.
Query March
Best Local Similarity 29.2%;
RESULT 855
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Local Similarity 29.2%;
                                  (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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JAN-2003.
(GETH) GENENTECH 1.
UNEY MATCH
BEST LOCAL SIMILARITY .
RESULT 854
ID ADA42646 stand?
DE HUMAN SECT?
PN US20030"
PD 20-"
                                                              Best Local Similarity
RESULT 849
                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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Best Local Similarity
US2003039969-A1.
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                                                     Query Match
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) 24-APR-2003.
1 (GETH ) GENENTECH INC.
Query Match 16.1%; Score 307.5; DB 7; Length 390;
16.1%; Score 307.5; DB 7; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                         Score 307.5; DB 7; Length 390; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                        16.1%; Score 307.5; DB 7; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                  Length 390;
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                                                ADABBS73 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003073213-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADB22268 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003087344-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                  Score 307.5; DB 7;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABO19865 standard; protein; 390 AA.
Human secreted/transmembrane protein PRO246.
US2003044902-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADB23593 standard; protein; 390 AA. Human PRO polypeptide SEQ ID NO 338 US2003077712-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human PRO polypeptide #169.
US200307711-A1..
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03-app.
                                                                                                                                                                                          ADB27335 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003022239-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADA66959 standard; protein; 390 AA
Human PRO polypeptide #169.
US2003068793-A1.
                                                                                                                                           16.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                  16.1%;
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Best Local Similarity 29.2%;
RESULT 865
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                    01-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                            (GETH ) GENENTECH INC
       Best Local Similarity RESULT 858
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Best Local Similarity
RESULT 866
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 863
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Best Local Similarity
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Best Local Similarity
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Best Local Similarity
                                                                                                                                                        Best Local Similarity
RESULT 859
                                                                                                                                                                                                                                                                                                                                                                                               30-JAN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-MAY-2003.
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                                                                                                           17-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                           Query Match
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ADB74701 standard, protein, 390 AA. Human secreted/transmembrane protein, #9.002003082542-A1. RESULT 876 Query Match 16.1%; Score 307.5; DB 7; Length 390; Best Local Similarity 29.2%; Pred. No. 4.4e-16; RESULT 868 Score 307.5; DB 7; Length 390; Pred. No. 4.4e-16; Score 307.5; DB 7; Length 390; Pred. No. 4.4e-16; Score 307.5; DB 7; Length 390; Pred. No. 4.4e-16; Score 307.5; DB 7; Length 390; Pred. No. 4.4e-16; DB 7; Length 390; Length 390; Score 307.5; DB 7; Length 390; Pred. No. 4.4e-16; Score 307.5; DB 7; Length 390; Pred. No. 4.4e-16; ADA92315 standard; protein; 390 AA. Novel human secreted and transmembrane protein PRO246. ADB38630 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246. US2003082766-A1. ADB38078 standard; protein; 390 AA. Novel human secreted and transmembrane protein PRO246. US2003087347-A1. ADB65550 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246. US2003082689-A1. Novel human secreted and transmembrane protein PRO246. US2003082764-A1. Score 307.5; DB 7; Pred. No. 4.4e-16; Best Local Similarity 29.2%; Pred. No. 4.4e-16; RESULT 867 Score 307.5; DB 7 Pred. No. 4.4e-16; ADB77565 standard; protein; 390 AA. Human secreted/transmembrane protein, #9 US2003077654-A1. ADB15378 standard; protein; 190 AA. Human PRO polypeptide #169. US2003087352-A1. Human PRO polypeptide #169. 01-Mav-20. Human PRO polypeptide #169.
US2003082698-A1. protein; 390 AA USZUOSU-01-MYZ-2003. (GETH) GENENTECH INC. MATCh --: larity 29.2%; US20030.... 24-APR-2003. (GETH) GENENTECH INC. March 16.1%; 16.1%; 16.1%; 16.1%; Match 16.1%; Local Similarity 29.2%; (GETH) GENENTECH INC. ry Match 16.1%; : Local Similarity 29.2%; Match 16.18; Local Similarity 29.28; 08-MAY-2003. (GETH) GENENTECH INC. 01-MAY-2003. (GETH) GENENTECH INC. 01-MAY-2003. (GETH) GENENTECH INC. 01-MAY-2003. (GETH) GENENTECH INC. (GETH) GENENTECH INC. (GETH) GENENTECH INC. Local Similarity Query Match Best Local Similarity Local Similarity Query Match Best Local Similarity ADB39463 standard; US2003082712-A1. 01-MAY-2003 01-MAY-2003 Query Match

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Query Match 16.1%; Score 307.5; DB 7; Length 390; Best Local Similarity 29.2%; Pred. No. 4.4e-16; RESULT 882
                                                      Score 307.5; DB 7; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                            DB 7; Length 390;
                                                                                                                                                                                                                                                                                                                                                             16.1%; Score 307.5; DB 7; Length 390; 29.2%; Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 390;
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                                                                                                                                                                                                                                                                                                                                                                                                               ADB77298 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003082696-Al.
                                                                                                             ADB47086 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003082687-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 7;
                                                                                                                                                                                                        16.1%; Score 307.5; DB 7 29.2%; Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                           ADB86693 standard; protein; 390 AA. Human PRO polypeptide #169. US2003082697-A1. GL-MAY-2003. (GFH.) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADB34455 standard; protein; 390 AA. Human PRO polypeptide SEQ ID NO 338 US2003077717-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADB35559 standard; protein; 390 AA. Human PRO polypeptide SEQ ID NO 338 US2003077719-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADB36111 standard; protein; 390 AA. Human PRO polypeptide SEQ ID NO 338 US2003077720-A1.
US2003002.
01-MAY-2003.
(GETH ) GENENTECH INC.
usrch 16.1%; Sc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16.1%;
29.2%;
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RESULT 884
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PA (GETH ) GENENTECH INC.

Query Match 16.1%;

Best Local Similarity 29.2%;

RESULT 885
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                           (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GETH ) GENENTECH INC.
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Best Local Similarity
                                                    Query Match
Best Local Similarity
RESULT 877
                                                                                                                                                                                                                        Best Local Similarity RESULT 878
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
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RESULT 879
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Human secreted/transmembrane protein, #9
                                                                                    Best Local Similarity RESULT 895
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                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZO-FED-ZOUD.
(GETH ) GENENTECH INC.
sty Match
16.1%; Score 307.5; DB 7; Length 390;
ar Tocal Similarity 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 307.5; DB 7; Length 390; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                          Score 307.5; DB 7; Length 390;
Pred. No. 4.4e-16;
                                                                                      Score 307.5; DB 7; Length 390; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                Score 307.5; DB 7; Length 390;
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 307.5; DB 7; Length 390; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 307.5; DB 7; Length 390;
Pred. No. 4.4e-16;
ADB46506 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246
US2003082692-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 307.5; DB 7;
Pred. No. 4.4e-16;
                                                                                                                                      ADC20347 standard, protein, 390 AA.
Human secreted/transmembrane protein, #9.
US2003059772-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADC34185 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003036094-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADC29240 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003049676-Al.
13-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADC40656 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003054400-A1.
                                                                                                                                                                                                                                                                                   ADC39547 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9
US2003059828-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                            ADC40061 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9
US2003059829-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADC18889 standard; protein; 390 AA.
Whaman secreted/transmembrane protein, #9.
US2003036061-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADC28771 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9
US2003049677-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ID ADC19313 standard; protein; 390 AA
                                    US2003-0
01-MPY-2003.
(GETH ) GENENTECH INC.
Match 17-rity 29.2%;
                                                                                                                                                                            US2003.2.27-MAK-2003.
(GETH ) GENENTECH INC.
(CTTA) CALL 16.1%;
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20-FEB-2003.
(GETH ) GENENTECH INC.
16.1%;
                                                                                                                                                                                                                                                                                                                        US2003.2.
27-MAR-2003.
(GETH ) GENENTECH INC.
Match 'Trity 29.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               US200305.2.
27-WAR-2003.
(GETH ) GENENTECH INC.
16.1%;
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(GETH ) GENENTECH INC.
:ry Match 16.1%;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 29.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
RESULT 888
ID ADC40061 standard; px
DE Human secreted/transm
PN US2003059829-A1.
PD 27-MAR-2003.
PA (GETH ) GENENTECH INC
                                                                                      Guery Match
Best Local Similarity
RESULT 886
ID ADC28347 standard; p
DE Human secreted/trans
PN US2003059772-A1.
PD 27-MAR-2003.
PA (GETH ) GENENTECH IN
                                                                                                                                                                                                                                  Query Match
Best Local Similarity
RESULT 887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
RESULT 890
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16.1%; Score 307.5; DB 7; Length 390; 29.2%; Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                  Score 307.5; DB 7; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 307.5; DB 7; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 16.1%; Score 307.5; DB 7; Length 390; Best Local Similarity 29.2%; Pred. No. 4.4e-16; RESULT 902
                                                                    Score 307.5; DB 7; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16.1%; Score 307.5; DB 7; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16.1%; Score 307.5; DB 7; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16.1%; Score 307.5; DB 7; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADC52912 standard; protein; 390 AA.
Novel human secreted and transmembrane protein Seg ID338.
US2003087365-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADC57266 standard; protein; 390 AA.
Novel human secreted and transmembrane protein Seg ID338.
US2003087366-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADCS0379 standard; protein; 390 AA. Novel human secreted and transmembrane protein PRO246.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADC71926 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003092107-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADC59905 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003092105-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADC60457 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003087367-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADC50932 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
                                                                                                                                                          Human secreted/transmembrane protein, #9. US2003073077-A1.
                                                                                                                                                                                                                                                                                                                                               Human secreted/transmembrane protein, #9.
US2003073079-A1.
                                                                                                                                     ADC33761 standard; protein; 390 AA
                                                                                                                                                                                                                                                                                                                         ADC12831 standard; protein; 390 AA
                                                                    16.1%;
                                                                                                                                                                                                                                                    16.1%;
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PA (GETH) GENENTECH INC.

Query Match 16.1%;

Best Local Similarity 29.2%;

RESULT 897
US2003054441-A1.
20-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                            17-APR-2003. (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 896
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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Length 390;

Length 390;

DB 7; Length 390;

7; Length 390;

Length 390;

DB 7; Length 390;

Length 390;

DB 7;

RESULT 911

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16.1%; Score 307.5; DB 7; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16.1%; Score 307.5; DB 7; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel human secreted and transmembrane protein PRO246. US2003087354-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADC80523 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003092103-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC79971 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003087358-A1.
08-MAY-2003.
                         Query Match 16.1%; Score 307.5; DB 7; Best Local Similarity 29.2%; Pred. No. 4.4e-16; RESULT 913
                                                                                                                                                                           Score 307.5; DB 7;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 307.5; DB 7;
Pred. No. 4.4e-16;
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Best Local Similarity 29.2%; Pred. No. 4.4e-16;
RESULT 918
                                                                                                                                                                                                                                                                                                                     Score 307.5; DB 7
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADD04838 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003104469-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human PRO polypeptide #169.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human PRO polypeptide #169.
1052003194771-A1.
                                                                                 ADC69574 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003194770-A1.
                                                                                                                                                                                                                                                                                                                                                                         ADD09992 standard; protein; 390 AA
                                                                                                                                                                                                                                390 AA
                                                                                                                                                                           16.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                16.1%;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 29.2%;
RESULT 917
                                                                                                                                                                                                                                                                                                                  Best Local Similarity 29.2%;
RESULT 915
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                                                                                                                                                                                                                             ADC48463 standard; protein;
Human PRO polypeptide #1,69.
US2003194773-A1.
                                                                                                                                                                                                                                                                                                                                                                                         Human PRO polypeptide #169
US2003194776-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                             16-OCT-2003.
(GETH ) GENENTECH INC.
                                                                                                                                          16-OCT-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                   (GETH ) GENENTECH INC.
                  GENENTECH INC.
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(GETH ) GENENTECH INC
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(GETH ) GENENTECH INC
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Best Local Similarity
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RESULT 919
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 08-MAY-2003
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                                                                                                                                                                             Query Match
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                                                   Score 307.5; DB 7; Length 390;
Pred. No. 4.4e-16;
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(GETH ) GENENTECH INC.
ry Match
- Toral Similarity 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                             Length 390;
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Pred. No. 4.4e-16;
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ry Match
t Local Similarity 29.2%; Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 390
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                                                                                                                                                                                                                                                                                                                                                                                         ADC53518 standard; protein; 390 AA.
Novel human secreted and transmembrane protein Seg ID338.
US2003087364-A1.
                                                                                                                                                                                                                                                                Novel human secreted and transmembrane protein Seq ID338 US2003087363-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AD 559041 standard; protein; 390 AA.
Novel human secreted and transmembrane protein Seg ID338
US2003087359-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADC58489 standard; protein; 390 AA.
Novel human secreted and transmembrane protein Seg ID338.
US2003087346-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADC55919 standard; protein; 390 AA.
Novel human secreted and transmembrane protein Seg ID338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADD03163 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003092104-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC90155 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003087348-A1.
                                                                                                                                                                                            Score 307.5; DB 7;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 307.5; DB 7;
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 307.5; DB 7;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADC12283 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003082541-A1.
                                                                                        Human PRO polypeptide #169.
US2003087362-A1.
                                                                                                                                                                                                                                                   protein; 390 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                           US2003uo,...
08-MAY-2003.
(GETH ) GENENTECH INC.
"Arch 16.1%; Sr
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-2003.
GENENTECH INC.
16.1%; S
                                                 16.1%;
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08-MAY-2003.
(GETH ) GENENTECH INC.
MAtch '12-ity 29.24;
                                                                                                                                                                                            16.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16.1%;
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Local Similarity 29.2%;
              08-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                         08-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GETH ) GENENTECH INC.
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                                                              Best Local Similarity
RESULT 904
                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                             ADC54557 standard;
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US2003087361-A1.
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                                                   Query Match
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16.1%; Score 307.5; DB 7; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16.1%; Score 307.5; DB 7; Length 390; 29.2%; Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16.1%; Score 307.5; DB 7; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                          16.1%; Score 307.5; DB 7; Length 390; 29.2%; Pred. No. 4.4e-16;
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Score 307.5; DB 7; Length 390. Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Abovel human secreted and transmembrane protein PRO246. US203194765-Al.
                                                                                                                                                                                            ADD54155 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
                                                                                                                                                                                                                                                                                                                                                                                                                         Score 307.5; DB 7;
Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                                                                                                   ADD92472 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003199030-A1.
                                                 ADD01973 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003203430-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADD91168 standard; protein; 390 AA. Human PRO polypeptide #169.
US2003199055-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADE22211 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003199056-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADD79435 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003203428-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADE41971 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003194772-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADE033982 standard; protein; 390 AA Human PRO polypeptide #169. US2003199057-A1.
 16.1%;
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Best Local Similarity 29.2%;
RESULT 937
                                                                                                                                                                                                                                                                                                                                                                                                                           16.18;
                                                                                                                                                                                                                                                                                                                                                                                                                                           29.28;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 938
         Best Local Similarity RESULT 931
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 936
                                                                                                                                                                                                                                                                                               Local Similarity
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                                                                                                                                                    Best Local Similarity RESULT 932
                                                                                                                                                                                                                                US2003203432-A1.
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                                                                                                                                          Query Match
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(GETH ) GENENTECH INC.
ery Match
if.if; Score 307.5; DB 7; Length 390;
if.if; Pred. No. 4.4e-16;
                  Score 307.5; DB 7; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                             DB 7; Length 390;
                                                                                                                                                                                                                                                                                                  Length 390;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADD41153 standard; protein; 390 AA. Novel human secreted and transmembrane protein PRO246. US2003203438-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DE ADD5384 standard; protein; 390 AA.
DE Novel human secreted and transmembrane protein PRO246.
PN . US2003203437-A1.
PD 30-0CT-2003.
PA (GETH.) GENENFECH INC.
                                                                                                                                                                                                                                                                                                    Score 307.5; DB 7;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                          16.1%; Score 307.5; DB 7; 29.2%; Pred. No. 4.4e-16;
                                                                 ADD09440 standard; protein; 390 AA.

Human PRO polypeptide #169.
US2003194775-A1.
(GETH) GENENTECH INC.

EXY MATCH

SEL Local Similarity 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 307.5; DB 7, Pred. No. 4.4e-16;
                                                                                                                                                                                                            ADD03844 standard; protein; 390 AA. Human secreted/transmembrane protein, #9. US2003104381-A1. 05-JUN-2003.
                                                                                                                                                                                                                                                                                                                                                  ADD03420 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
22003108983-A1.
12-JUN-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human PRO polypeptide #169.
30-Orm-on.
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16-0003194769-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADD53032 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003194792-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein; 390 AA
                                                                                                                                                                                                                                                 JS20031.
05-77N-2003.
(GETH ) GENENTECH INC.
16.1%;
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30-007-2003.
(GETH ) GENENTECH INC.
16.1%;
                  16.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16.1%;
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29.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human PRO polypeptide #169.
US2003194779-A1.
16-OCT-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-OCT-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity RESULT 930
                                                                                                                                                               Best Local Similarity
RESULT 923
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Best Local Similarity
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Best Local Similarity
RESULT 927
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                                Local Similarity
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                    Query Match
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Length 390;

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Query Match
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                                                                                                                      DB 7; Length 390;
                                                                                                                                                                                                                                                              Score 307.5; DB 7; Length 390;
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                                                                             ADE33383 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003194767-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADE33935 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003194791-A1.
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Pred. No. 4.4e-16;
29.2%; Pred. No. 4.4e-16;
                                                                                                                      Score 307.5; DB 7
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Human secreted/transmembrane protein, #90
US2003077583-A1.
                                                                                                                                                                 ADD91920 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003199053-A1.
                              ADE17788 standard; protein; 390 AA. Human PRO polypeptide #169. US2003199023-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human PRO polypeptide #169.
US2003207417-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADE19444 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003199025-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADD93024 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003194768-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein; 390 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2003123
23-007-2003.
(GETH ) GENENTECH INC.
16.1%;
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Local Similarity 29.2%;
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Best Local Similarity 29.2%;
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Local Similarity 29.2%;
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Best Local Similarity 29.2%;
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Best Local Similarity 29.2%;
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Human PRO polypeptide #169.
US2003199026-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-OCT-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                            23-OCT-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                  16-OCT-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                     (GETH ) GENENTECH INC.
                                                                                                                                      Local Similarity
Best Local Similarity
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Best Local Similarity
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Best Local Similarity
                                                                                     23-OCT-2003
                                                                                                                      Query Match
                                                                                                                                                                                                                                                              Query Match
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                                                                                                                                                        RESULT 941
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unery Match 16.1%; Score 307.5; DB 7; Length 390; Best Local Similarity 29.2%; Pred. No. 4.4e-16; SULT 954
                                                                                    Length 390;
                                                                                                                                                                                                                DB 7; Length 390;
                                                                                                                                                                                                                                                                                                                                             Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 390;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADE32831 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003194766-A1.
                                                                                    DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 7;
                                                                                  Score 307.5; DB 7
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                                                                                                             Score 307.5; DB 7
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
ADE43088 standard; protein; 390 AA.
Human PRO polypeptide #169.
US203139033-A1.
23-OCT-2003.
(GETH ) GENENTECH INC.
Cuery Match
16.1%; Score 30.
3111.7 96.0
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0S2003207418-A1.
                                                                                                                                                                                                                                                          ADE22763 standard; protein; 390 AA. Human PRO polypeptide #169. US2003199064-A1.
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Human PRO polypeptide #169.
US2003199031-A1.
                                                                                                                                 protein; 390 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADE42523 standard; protein; 390 AA
Human PRO polypeptide #169.
US2003199032-A1.
                                                                                                                                                                                                                                                                                                                                           16.1%;
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PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 29.2%;
RESULT 957
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Best Local Similarity 29.2%;
RESULT 953
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RESULT 955
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PA (GETH ) GENENTECH INC.
Query Match 16.1%;
Best Local Similarity 29.2%;
RESULT 958
                                                                                                                                               Human PRO polypeptide #169.
US2003199059-A1.
23-OCT-2003.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                             23-OCT-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
SULT 956
                                                                                                                                                                                                                            Best Local Similarity
RESULT 951
                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 952
                                                                                            Best Local Similarity RESULT 950
                                                                                                                                 ADD95877 standard;
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Best Local Similarity
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                                                                               Best Local Similarity
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ID ADJ26202 standard;
                                                 (GETH ) GENENTECH
                                                                                                                                                                                                                                                                                                 20-MAR-2003
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RESULT 974
                                                                               Score 307.5; DB 7; Length 390; Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                          ADG21488 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207355-A1.
                                                                                                                                                                                                                                                                                                                                                                                         ADG21129 standard, protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207384-A1.
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Pred. No. 4.4e-16;
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Human secreted/transmembrane polypeptide PRO246
US2003170721-A1.
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US2003207373-A1.
ADE04650 standard, protein, 390 AA.
Human PRO polypeptide #169.
US2003199034-A1.
33-OCT-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADF97464 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003207370-A1.
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Human PRO polypeptide #169.
US2003207372-A1.
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UGZCH 2003.
(GFCH ) GENENTECH INC.
TWATCh ''larity 29.2*;
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06-NOV-2003.
(GETH ) GENENTECH INC.
16.1%;
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06-NOV-2003.
(GETH ) GENENTECH INC.
MATCh '''1arity 29.2%;
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06-NOV-2003.
(GETH ) GENENTECH INC.
16.1%;
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RESULT 963
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Best Local Similarity 29.2%;
RESULT 967
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Local Similarity 29.2%;
                                                                                                                                              Human PRO polypeptide #169
US2003194777-A1.
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Best Local Similarity
RESULT 960
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                                                                                              Best Local Similarity
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                                                                                                                                ADE92779 standard;
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16.1%; Score 307.5; DB 7; Length 390; 29.2%; Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
                                                                                                 Length 390;
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Novel human secreted and transmembrane protein PRO246.
US2003207381-A1.
06-NOV-2003.
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Novel human secreted and transmembrane protein PRO246.
US2003207385-A1.
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Novel human secreted and transmembrane protein PRO246.
US2003207388-A1.
                                                                                                                                                                ADM55820 standard; protein; 390 AA.

Novel human secreted and transmembrane protein PRO246.
US2003207379-A1.
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Novel human secreted and transmembrane protein PRO246.
US2003207386-A1.
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Novel human secreted and transmembrane protein PRO246.
US2003207387-A1.
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Novel human secreted and transmembrane protein PRO246.
US2003207377-A1.
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Novel human secreted and transmembrane protein PRO246.
                                                                                                 DB 7;
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 29.2%; Pred. No. 4.4e-16;
                                                                                              16.1%; Score 307.5; DB 7 29.2%; Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                                                                                                                       ADI37934 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003054352-A1.
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29.2%;
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Best Local Similarity 29.2%;
RESULT 970
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Length 390;

Length 390;

Length 390;

Length 390;

8

Length 390;

us-10-002-967a-59.rag.spdi

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16.1%; Score 307.5; DB 8; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                  Score 307.5; DB 8;
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Human secreted/transmembrane polypeptide PRO246.
US2003104558-A1.
05-UNN-2003.
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Pred. No. 4.4e-16;
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                                                                                                                           ADE79541 standard; protein; 390 AA.
Human secreted/transmembrane protein,
US2003130489-A1.
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US2003211571-A1.
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Human PRO polypeptide #169.
US2003092108-A1.
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US2003092110-A1.
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Human PRO polypeptide #169.
US2003203439-A1.
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Human PRO polypeptide #169.
US2003199062-A1.
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(GETH ) GENENTECH INC.
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15-WAY-2003.
(GETH ) GENENTECH INC.
Match 'Towity 29.2%;
                                                                                                                                                                                                                                                            16.1%;
                     GETH GENENTECH INC.
Query Match
16.1%;
Best Local Similarity 29.2%;
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Best Local Similarity 29.2%;
RESULT 989
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                                                                                                                                                                                                   (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 991
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Best Local Similarity
RESULT 987
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Best Local Similarity
  30-OCT-2003
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                                                                           16.1%; Score 307.5; DB 7; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                    Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 307.5; DB 7; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ), 08-MAY-2003.
A (GETH) GENENTECH INC.
Query Match
Best Local Similarity 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 307.5; DB 8; Length 390;
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 390;
                                                                                                                              AUN15917 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US203087353-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADNI5365 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003087356-A1.
                                                                                                                                                                                                                                                                                                                                                            ADNI6546 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003087385-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADN14813 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003087357-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADC81075 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003092115-A1.
                                                                                                                                                                                                                                                                                       DB 7;
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Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                       Score 307.5; DB 7
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADE79117 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003135025-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human PRO polypeptide #169.
. US200320340-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein; 390 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          390 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US200300...
08-MAY-2003.
(GETH ) GENEWTECH INC.
16.1%;
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( (GETH ) GENENTECH INC.

Query Match 16.1%;

Best Local Similarity 29.2%;
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Best Local Similarity 29.2%;
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Local Similarity 29.2%;
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Local Similarity 29.2%;
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Best Local Similarity 29.2%;
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US2003100087-A1.
                                                                                                                                                                                                                              08-MAY-2003.
(GETH ) GENENTECH INC.
                        08-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                   Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADD76523 standard;
US2003087355-A1.
                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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DB 8; Length 390;

Length 390;

DB 8;

RESULT

Length 390;

8;

Query Match Best Local Similarity 29.2%; F RESULT 1004 ID ADE99455 standard; protein; 39 DE Human secreted/transmembrane p DI 3-007-2003. PA (GETH) GENENTECH INC. Query Match Best Local Similarity 29.2%; F RESULT 1005 ID ADE92227 standard; protein; 39 DE Novel human secreted and trans PH 05200199051-A1.	255 25	DE ADE91675 standard; protein; 39 DE Novel human secreted and trans PN US2003199058-A1. PD 23-OCT-2003. PA (GETH) GENENTECH INC. Query Match Best Local Similarity 29.2%; E RESULT 1008 ID ADE98852 standard; protein; 39 DE Human secreted/transmembrane F PN US2003211568-A1. PD 13-NOV-2003.	3 5 5	PA (WILL) WILLIAMS P W. PA (WILL) WILLIAMS P W. PA (WOOD/) WOOD W I. Query Match Bast Local Similarity 29.2%; E RESULT 1010 ID ADF73716 standard; protein; 35 DE Human secreted/transmembrane F PN US2003180312-A1. PD 25-SEP-2003. PA (GETH) GENENTECH INC. Query Match Query Match 16.1%; S	αgi Ri
7.5; DB 8; Length 390; 4.4e-16; #9. 7.5; DB 8; Length 390;	.5; DB 8; Length 390;	, Q 104,	5; DB 8; Length 390; 4.4e-16; 5; DB 8; Length 390; 4.4e-16;	5; DB 8; Length 390; 4.4e-16;	5; DB 8; Length 390; 4.4e-16;
Score 30' Pred. No 390 AA. Sprotein, Score 30' Pred. No	Score 307. Pred. No. 90 AA. Score 307.	90 AA. Score 307 Pred. No.	INC. 16.1%; Score 307.5; 29.2%; Pred. No. 4. protein; 390 AA. ide #169. INC. 16.1%; Score 307.5; 29.2%; Pred. No. 4.	protein; 390 AA. ide #169. INC. 16.1%; Score 307.5; 29.2%; Pred. No. 4. protein; 390 AA.	INC. 16.1%; Score 307.5; 29.2%; Pred. No. 4. protein; 390 AA. ide #169.
DA (GETH) GENENTECH INC. Query Match Best Local Similarity 29.2%; Pred. No RESULT 995. DB Human secreted/transmembrane protein, DN 07-AUG-2003. PA (GETH) GENENTECH INC. Query Match Best Local Similarity 29.2%; Pred. No RESULT 996 ID ADE18340 standard; protein; 390 AA. DE Human PRO polypeptide #169. PA (GETH) GENENTECH INC. Query Match RESULT 996 ID ADE18340 standard; protein; 390 AA. DE Human PRO polypeptide #169.	,ă, ă, ă, ă,	RESULT 998 ID ADE99306 standard; protein; 390 AA. DE Human secreted/transmembrane protein, PN US2003211576-A1. PD 13-NOV-2003. PA (GETH) GENENTECH INC. GUETY MATCh Best Local Similarity 29.2%; Pred. NO RESULT 999 ID ADE94669 standard; protein; 390 AA. DE Human PRO Polypeptide #169.	igg ggg	KESULI 1001 ID ADE95521 standard; protein; DE Human PRO polypeptide #169. PN US2003199052-A1. PP 23-OCT-2003. PA (GETH) GENENTECH INC. QUETY MATCh Best Local Similarity 29.2%; RESULT 1002 ID ADE93331 standard; protein; DE Human PRO polypeptide #169. PN US2003199060-A1.	PA (GETH) GENENTECH INC. Query Match Best Local Similarity 29.2%; RESULT 1003 ID ADP34912 standard; protein; DE Human PRO polypeptide #169. PD US2003199029-A1. PA (GETH) GENENTECH INC.

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Score 307.5; DB 8; Length 390; Pred. No. 4.4e-16;
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Score 307.5; DB 8; Length 390; Pred. No. 4.4e-16;
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                                                                                                                                                                        390 AA.
nsmembrane protein PRO246.
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nsmembrane protein PRO246.
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protein, #9.
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: protein, #9.
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protein, #9.
                                             390 AA.
protein, #9.
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PA (GETH) GENENTECH INC.

Query Match

16.1%; Score 307.5; DB 8; Length 390;

Best Local Similarity 29.2%; Pred. No. 4.4e-16;

RESULT 1015
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     / Match 16.1%; Score 307.5; DB 8; Length 390; Local Similarity 29.2%; Pred. No. 4.4e-16;
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                                                                                                Length 390;
                                                                                                                                                                                                                                                              16.1%; Score 307.5; DB 8; Length 390; 29.2%; Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
ADG22040 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207360-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADG24233 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246
                                                                                                Score 307.5; DB 8;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 307.5; DB 8;
Pred. No. 4.4e-16;
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US2003208055-Al.
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Human PRO polypeptide #169.
US2003207376-A1.
                                                                                                                                                                                                                                                                                                                  ADF98016 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003207422-A1.
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Human PRO polypeptide #169.
US2003207351-A1.
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Human PRO polypeptide #169.
US2003207359-A1.
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Human PRO polypeptide #169.
US2003207375-A1.
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                                                        PD 06-NOV-2003.

PA (GETH) GENENTECH INC.

QUELY MATCh 16.1%;

BEST LOCAL Similarity 29.2%;

RESULT 1013
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Best Local Similarity 29.2%;
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Local Similarity 29.2%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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06-NOV-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                 Best Local Similarity
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Best Local Similarity 29.2%; Pred. No. 4.4e-16; RESULT 1025
ID ADG15514 standard; protein; 390 AA.
BN US2003119885-A1.
                                                                                                                                                                                                                                                                                                                                                             PD 06-NOV-2003.

PA (GETH ) GENENTECH INC.

Query Match

Best Local Similarity 29.2%; Pred. No. 4.4e-16;

RESULT 10.24

ID ADG08344 standard; protein; 390 AA.

DE Novel human secreted and transmembrane protein PRO246.

PN US2003207424-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ouery Match 16.1%; Score 307.5; DB 8; Length 390; Best Local Similarity 29.2%; Pred. No. 4.4e-16; RESULT 1027
                                                                                                                                                                                                                                           Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16.1%; Score 307.5; DB 8; Length 390; 29.2%; Pred. No. 4.4e-16;
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Best Local Similarity 29.2%; Pred. No. 4.4e-16;
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                                                                              Length 390;
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Novel human secreted and transmembrane protein PRO246.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADG23681 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246
US2003207389-A1.
                                                                                                                                                                                                                                             DB 8;
                                                                              Score 307.5; DB 8;
Pred. No. 4.4e-16;
                                                                                                                                                                                                   PD 04-SEP-2003.

PA (GETH ) GENENTECH INC.

QUENT BEST LCCAL Similarity 29.2%; Pred. No. 4.4e-16; RESULT 1023

ID ADG13287 standard; protein; 390 AA.

BY US2003207357-AI.
                                                                                                                                     ADF71292 standard; protein; 390 AA.
Human secreted/transmembrane protein,
US2003166051-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human PRO polypeptide #169.
US2003207423:Al.
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Human PRO polypeptide #169.
US2003207371-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADG06097 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2003207374-A1.
                                                J6-NOV-2-0---
(GETH ) GENENTECH INC.
ry Match 16.1%;
Human PRO polypeptide #169.
US2003207425-A1.
06-NOV-2003.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                Query Match
Best Local Similarity
RESULT 1022
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
RESULT 1030
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Best Local Similarity
RESULT 1028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
RESULT 1026
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1029
ID ADG03970 standard;
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(GETH ) GEN
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DB 8; Length 390;

Length 390;

28

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Score 307.5; DB 8; Length 390; Pred. No. 4.4e-16;
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16.1%; Score 307.5; DB 8; Length 390;
Best Local Similarity 29.2%; Pred. No. 4.4e-16;
RESULT 1046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16.1%; Score 307.5; DB 8; Length 390; 29.2%; Pred. No. 4.4e-16;
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Best Local Similarity 29.2%; Pred. No. 4.4e-16;
RESULT 1042
ID ADG70893 standard; protein; 390 AA.
DE Novel human secreted and transmembrane protein PRO246.
PN US2003207420-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MDG57975 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207363-AI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A. ADG55559 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207415-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADG71445 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207421-A1.
                                                                                          ADGS5767 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207365-A1.
                                                                                                                                                                                                                                              ADGS6527 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207368-A1.
                                     16.1%; Score 307.5; DB 8 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 307.5; DB 8 Pred. No. 4.4e-16;
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PA (GETH ) GENENTECH INC.

Query Match 16.1%; Score 307.5; DB 8

Best Local Similarity 29.2%; Pred. No. 4.4e-16;

RESULT 1045
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human secreted/transmembrane protein, #9.
US2003027146-A1.
06-FEB-2003.
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082003207805-Al.
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24-App.
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ADG92562 standard; protein; 390 AA
                                                                                                                                                                                         16.1%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                     (GETH ) GENENTECH INC
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                                         Query Match
Best Local Similarity
                                                                                                                                                                                                    Best Local Similarity
RESULT 1041
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Best Local Similarity
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   06-NOV-2003
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PA (GETH) GENENTECH INC.

Query Match

Best Local Similarity 29.2%; Pred. No. 4.4e-16;

RESULT 1037
                                                         Score 307.5; DB 8; Length 390; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 307.5; DB 8; Length 390;
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Novel human secreted and transmembrane protein PRO246.
US2003207362-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADG56871 standard, protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207364-A1.
                                                                                                              ADG07168 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207350-A1.
                                                                                                                                                                                                                                              7T 1032
ADG07720 standard; protein; 390 AA.
ADG07720 standard; protein; 390 AA.
BYS00320120036-A1.
GS003207356-A1.
GGFH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                            Novel human secreted and transmembrane protein PRO246. US2003194778-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADG66879 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207390-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADG61983 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207428-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 307.5; DB 8;
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Human secreted/transmembrane protein, #9.
US2003027145-A1.
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06-NOV-2003.
(GETH ) GENENTECH INC...
Match 15.1%;
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06-000-2003.
(GETH ) GENENTECH INC.
16.1%;
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(GETH) GENENTECH INC.
If Match
If I similarity 29.2%;
                                                         16.1%;
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ry Match 16.1%;
t Local Similarity 29.2%;
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Local Similarity 29.2%;
                                       GENENTECH INC.
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                                                                          Local Similarity
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US2003207427-A1.
                     06-NOV-2003
(GETH ) GEN
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DB 8; Length 390;

DB 8; Length 390;

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Best Local Similarity 29.2%; Pred. No. 4.4e-16;
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(GETH ) GENENTECH INC.
16.1%; Score 307.5; DB 8; Length 390;
16.1%; Score 307.5; DB 8; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                Length 390;
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Local Similarity 29.2%; Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Novel human secreted and transmembrane protein PRO246.
US2003207416-A1.
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Novel human secreted and transmembrane protein PRO246.
US2003207366-A1.
                                                                                                                                                                                                                        ADH11961 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207419-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADH12585 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207378-A1.
                                                                                                                                                                                                                                                                                                                                                                                     ADG$2383 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207414-A1.
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Novel human secreted and transmembrane protein PRO246.
                                                                                                                                                              Score 307.5; DB 8;
Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
                                                       ADG63640 standard; protein; 390 AA.
Human secreted/transmembrane polypeptide PRO246.
US2003180796-A1.
25-SEP-2003.
                                                                                                                                                                                                                                                                                                                             Query Match 16.1%; Score 307.5; DB 8
Best Local Similarity 29.2%; Pred. No. 4.4e-16;
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Human PRO polypeptide #169.
US2003194793-A1.
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                                                                                                                                                            16.1%;
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Local Similarity 29.2%;
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Best Local Similarity 29.2%;
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1 (GETH) GENENTECH INC.

16-DETY MATCH 16.1%;

Best Local Similarity 29.2%;
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US2003022331-Al.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                 Local Similarity
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Best Local Similarity
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Query Match
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06-NOV-2003.
(GETH) GENENTECH INC.
ery Match 16.1%; Score 307.5; DB 8; Length 390;
                                                                                                                                                                                                                                               Length 390;
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ADG54663 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207367-A1.
                                                                                                                                                    ADGS9703 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
                                                                                                                                                                                                                                           16.1%; Score 307.5; DB 8; 29.2%; Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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Pred. No. 4.4e-16;
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                                                                                                                                                                                                                                                                                                                    Human secreted/transmembrane protein, #9. US2004005553-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADH07206 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2004006211-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human secreted/transmembrane protein, #9.
US2003215904-A1.
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Human secreted/transmembrane protein, #9.
US2004005665-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADI18521 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003152999-A1.
                                                                                                                                                                                                                                                                                                    ADH20351 standard; protein; 390 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADH59751 standard; protein; 390 AA
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Human PRO polypeptide #169.
US2003207361-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (GURN/) GURNEY A L.
(MATH/) MATHER J P.
(WILL/) WILLIAMS P M.
(WOOD/) WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GURNEY A L.
MATHER J P.
WILLIAMS P M.
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RESULT 1065
                                                                                                         Best Local Similarity RESULT 1059
                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                           Best Local Similarity
RESULT 1060
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GODDARD A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GODDARD A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WOOD W I.
                                                                                                                                                                                         US2003207369-A1
                                                                                                                                                                                                          06-NOV-200
                                                                                                                                                                                                                                                                                                                                                               08-JAN-200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-NOV-2003
                                                                                               Ouery Match
                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 1061
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DESNOYERS L.

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(DESN/)
         Score 307.5; DB 8; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                 Score 307.5; DB 8; Length 390; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PD 06-NOV-2003.

PA (GETH) GENENTECH INC.
Query Match 16.1%; Score 307.5; DB 8; Length 390;
Best Local Similarity 29.2%; Pred. No. 4.4e-16;
RESULT 1071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-JAN 2004.
(GETH ) GENENTECH INC.
ery Match 16.1%; Score 307.5; DB 8; Length 390;
                                          DB 8; Length 390;
                                                                                                                                                                                                                                                                                                                           Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 307.5; DB 8; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 307.5; DB 8; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 8; Length 390;
                                                                                                                                                                                                                                                                                                                                                                   ADG09870 standard; protein; 390 AA.

Novel human secreted and transmembrane protein PRO246.
US2004009548-AI.
15-JAN-2004.
(GETH.) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADII5341 standard, protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207382-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADI14673 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207383-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 307.5; DB 8;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 29.2%; Pred. No. 4.4e-16;
RESULT 1073
                                        Score 307.5; DB 8;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human secreted/transmembrane protein, #9. US20610-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADH60411 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US200402331-A1.
05-FBB-2004.
                                                                                         ADI65241 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003148419-A1.
                                                                                                                                                                                                                                   ADI37504 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9
US2003096340-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADI65668 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9
                                                                                                                            US200312-2
07-ADC-2003.
(GETH ) GENENTECH INC.
16.1%;
                                                                                                                                                                                                                                                                             USZOWZZ
22-MAZ-2003.
(GETH ) GENENTECH INC.
MATCh 'Intity 29.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USZUCZ-2003.
09-007-2003.
(GETH ) GENENTECH INC.
MATCh -- 13-rity 29.28;
PD 14-AUG-2003.

PA (GETH ) GENENTECH INC.

QUETY MATCh 16.1%;

Best Local Similarity 29.2%;

RESULT 1066
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 29.2%;
RESULT 1069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 16.1%;
Best Local Similarity 29.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ID AD165668 standard; pro
DE Human secreted/transmer
PN US2003148371-A1.
PD 07-AUG-2003.
PA (GETH) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
RESULT 1068
ID ADG09870 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
RESULT 1070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 1072
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-NOV-2003
                                                                                                                                                                                 Query Match
Best Local Si
RESULT 1067
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                                                                                                                                                                                                                                                                                                                            Query Match
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Query Match 16.1%; Score 307.5; DB 8; Length 390; Best Local Similarity 29.2%; Pred. No. 4.4e-16; RESULT 1079
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 307.5; DB 8; Length 390; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16.1%; Score 307.5; DB 8; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 307.5; DB 8; Length 390; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 390;
                                                                                                                16.1%; Score 307.5; DB 8; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16.1%; Score 307.5; DB 8; Length 390; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                             DB 8; Length 390,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AD165549 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2004039164-A1.
                                                                                                                                                                          ADI18268 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207349-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 307.5; DB 8;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                    16.1%; Score 307.5; DB 8; 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                             16.1%; Score 307.5; DB 8 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADM29752 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003190611-A1.
                                                                                                                                                                                                                                                                                                                                    ADJ99468 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003187238-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADLO8661 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003186358-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 1078

ID ADM25006 standard; protein; 390 AA.

DE Human secreted/transmembrane protein, #9.

PN US2003096233-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human PRO polypeptide #169. US200403835-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADJ77444 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2004038336-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16.1%; 29.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 16.1%;
Best Local Similarity 29.2%;
RESULT 1081
(GODD/) GODDARD A.
(GODO/) GODOWSKI P J.
(GURN/) GURBY A L.
(MATH/) MATHER J P.
(WILL/) WILLIAMS P M.
(WOOD/) WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-OCT-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                     06-NOV-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-FEB-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                    (GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 1077
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
SULT 1080
                                                                                                                                                                                                                                                                                         Best Local Similarity RESULT 1076
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                                                                                                                                   Best Local Similarity RESULT 1075
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                                                                                                                                                                                                                                                                                                                                                                                                    02-OCT-2003
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                                                                                                                     Query Match
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ADS74474 standard; protein; 390 AA.
Human secreted/transmembrane protein #9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADH62551 standard; protein; 268 AA.
Human A33 antigenic protein fragment #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 29.2%; Pred. RESULT 1094
ID AAY08073 standard; protein; 268 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16.0%;
                                    PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 29.2%;
RESULT 1090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16.0%;
    Human PRO polypeptide #169.
US2003077659-A1.
                                                                                                                                                                                                                                                                 14-AUG-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GERBER H.
GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (WILL/) WILLIAMS P M. (WOOD/) WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                             BOTSTEIN D.
DESNOYERS L.
EATON D L.
                                                                                                                                                                                                                                                                                                                                                                                               ASHK/) ASHKENAZI A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILVAROFF E. FONG S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STEWART T A.
TUMAS D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity RESULT 1095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KLJAVIN I J
MATHER J P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
RESULT 1093
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PAONI N F.
ROY M A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GODOWSKI P
GRIMALDI C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GURNEY A L
                                                                                                                                                                                                                                                                                                                                                                                                                                                        FERRARA N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human A33 protein.
WO9914241-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GODDARD A
                                                                                                                                                                                                                                                                                                                                                                    JS2004185531-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (REED/) REED R. (ZHOU/) ZHOU Z.
                                  24-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GERR/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PAON/)
                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 29.2%; Pred. No. 4.4e-16; RESULT 1087

ID ADRI0926 standard; protein; 390 AA.

DB Human secreted/transmembrane protein, #9.

PN US2004137561-A1.
                                                                                                                                                                                                                                                                                                                         Score 307.5; DB 8; Length 390; Pred. No. 4.4e-16;
                                                                                                                                                                                                      Score 307.5; DB 8; Length 390; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 16.1%; Score 307.5; DB 8; Length 390; Local Similarity 29.2%; Pred. No. 4.4e-16;
                                                                                       Score 307.5; DB 8; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 307.5; DB 8; Length 390;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human secreted/transmembrane protein, #9.
US20041417017-A1.
                                                                                                                            ADM42426 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2004058424-A1.
                                                                                                                                                                                                                                          AD006074 standard; protein; 390 AA. Human PRO polypeptide #8. US6686451-B1.
               ADM27702 standard; protein; 390 AA.
Human PRO polypeptide #169.
US2004048333-Al.
                                                                                                                                                                                                                                                                                                                                                                ADM28288 standard, protein, 390 AA.
Human PRO polypeptide #169.
US2004077064-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADI95770 standard; protein; 390 AA
                                          US20040-1
11-MAR-2004.
(GETH ) GENENTECH INC.
Warch 16.1%; S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16.1%;
29.2%;
                                                                                                                                                                                                     Query Match 16.1%;
Best Local Similarity 29.2%;
                                                                                                                                                                                                                                                                                                                       / Match 16.1%;
Local Similarity 29.2%;
                                                                                                                                                                       25-MAR-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                          03-FEB-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                          (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-JUL-2004.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BOTSTEIN D.
DESNOYERS L.
EATON D L.
FERRARA N.
FILVAROFF E.
FONG S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PAN J.
PAONI N F.
ROY M A.
STEWART T A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TUMAS D.
WILLIAMS P M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GRIMALDI C J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ASHKENAZI A.
                                                                                              Best Local Similarity RESULT 1084
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 1089
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HILLAN K J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KLJAVIN I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GERBER H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GURNEY A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GAO W.
                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ASHK/)
(BOTS/)
RESULT 1083
ID ADM27700
DE Human Pi
PN US20040
PD 11-MAR-PA (GETH )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (DESN/)
(EATO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MOOD/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PANJ/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  KLJA/
MATH/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GAOW/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query
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Score 307.5; DB 8; Length 390; Pred. No. 4.4e-16;
                                                                                                                                                                                               Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                          DB 8; Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 304.5; DB 2; Length 268; Pred. No. 4.6e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 305.5; DB 6; Length 390;
Pred. No. 6.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABO53064 standard, protein, 390 AA.
Human putative spliceosome associated protein (SAP) #41
US2003068803-A1.
                                                                 AD196322 standard; protein; 390 AA.
Novel human secreted and transmembrane protein PRO246.
US2003207354-A1.
                                                                                                                                                PD 06-NOV-2003.

PA (GETH) GENENTECH INC.

Query Match 16.1%; Score 307.5; DB 8;

Best Local Similarity 29.2%; Pred. No. 4.4e-16;

RESULT 1091
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 307.5; DB 8;
Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 16.1%; Score 307.5; DB 8
Best Local Similarity 29.2%; Pred. No. 4.4e-16;
                                                                                                                                                                                                                                                                      ADT03511 standard; protein; 390 AA.
Human secreted/transmembrane protein, #9.
US2003152922-A1.
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity RESULT 1108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 1111
                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity RESULT 1106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                       (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-MAY-2004
                                                                  30-MAR-2000
                                                                                                                                                                                                                                                      Ouery Match
                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                             GOS-JUN-1999.
(GETH ) GENENTECH INC.
ery Match 16.0%; Score 304.5; DB 2; Length 270;
ery Match 31.9%; Pred. No. 4.6e-16;
                                                                                                                                                                                                                                                                                                                                                                                       03-JUN-1999.
(IGETH ) GENENTECH INC.
16.0%; Score 304.5; DB 2; Length 273; t Local Similarity 31.9%; Pred. No. 4.7e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 303.5; DB 4; Length 321;
Pred. No. 7e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15.9%; Score 303.5; DB 8; Length 321; 30.2%; Pred. No. 7e-16;
                                                                                                                                                    DB 7; Length 268;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 304.5; DB 2; Length 273; Pred. No. 4.7e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 304.5; DB 7; Length 273; Pred. No. 4.7e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 303.5; DB 4; Length 321;
Pred. No. 7e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human viral receptor protein homologue, SEQ ID NO:2307. WO200157188-A2.
                                                                                                                                                  16.0%; Score 304.5; DB 7, 31.9%; Pred. No. 4.6e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human therapeutic contig protein; 321 AA.
Human therapeutic contig protein - SEQ ID 2055.
23-SEP-2004.
(NUVELO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human A33 antigenic protein; 273 AA. Human A33 antigenic protein fragment #2. 11-SRD-2003171568-A1.
                                                                                                                                                                                                                                                                                                                                    AAY23329 standard; protein; 273 AA.
An A33 related antigen sequence.
WO9927098-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similar.
RESULT 1098
ID AAY08075 standard, protein; 273 AA.
DE Human A33 protein fragment #2.
PN W09914241-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAM40551 standard; protein; 321 AA.
Human polypeptide SEQ ID NO 5482.
WO200153312-A1.
                                                                                                                                                                                                     AAY23327 standard; protein; 270 AA.
                                                                                                                                                                                                                   An A33 related antigen sequence. WO9927098-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 16.0%;
Local Similarity 31.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-1999.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ASHK) ASHKENAZI A.
(PONG) FONG S.
(GOND) GODDARD A.
(GURN) GURNEY A L.
(TUMA/) TUMAS D.
(WODD/) WOOD W I.
                                                                                  (GURN/) GURNEY A L.
(NAPI/) NAPIER M A.
(TUMA/) TUMAS D.
(WOOD/) WOOD W I.
                          ASHKENAZI A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 1101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                FONG S.
GODDARD A.
                                                                                                                                                                  Best Local Similarity RESULT 1096
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-JUL-2001.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HYSE-) HYSEQ INC.
US2003171568-A1.
               11-SEP-2003.
(ASHK/) ASHKO
(FONG/) FONG
(GODD/) GODDI
(GURN/) GURNI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                       Query Match
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AABO8903 standard; protein; 327 AA.
Human secreted protein sequence encoded by gene 13 SEQ ID NO:60.
WO200017222-A1.
                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 29.9%; Pred. No. 7.6e-16;
RESULT 1105
ID ADE08316 standard; protein; 538 AA.

BN WO2003054152-A2.

PD 03-JUL-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          y Match 15.7%; Score 299.5; DB 4; Length 390; Local Similarity 28.9%; Pred. No. 1.9e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 7; Length 406;
                                                                                      (HUMA-) HUMAN GENOME SCI INC.
ry Match
t Local Similarity 29.9%; Pred. No. 7.2e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   D AAY67312 standard; protein; 387 AA.

E Human secreted protein AJ26_3 amino acid sequence.

N W09950020-A1.

D 25-NOV-1999.

A (GEMY ) GENETICS INST INC.

OUERY MAtch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 387;
                                                                                                                                                                                                                                                                                                                                Length 317;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 348;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 370;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 538;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AD043535 standard; protein; 348 AA.
Amino acid sequence of an additional human A34 clone
WO2004037999-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAE06610 standard; protein; 390 AA.
Human protein having hydrophobic domain, HP10801
WO200149728-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-JUN-1999.
(GEMY ) GENETICS INST INC.
15.5%; Score 295; DB 2;
27 Match
26.6%; Pred. No. 4.4e-15;
                                                                                                                                                                                                                                                                                                                                     15.9%; Score 303; DB 7; 29.9%; Pred. No. 7.6e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15.8%; Score 302; DB 7; 25.9%; Pred. No. 1.9e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PD 13-MAY-2004.
PA (GETH ) GENENTECH INC.
Query Match 15.8; Score 295; DB 8;
Best Local Similarity 26.68; Pred. No. 4.2e-15;
RESULT 1110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15.5%; Score 296.5; DB 7 27.1%; Pred. No. 3.6e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (LUDM-) LUDWIG INST CANCER RES.

ry Match
15.5%; Score 295; DB 8;

Local Similarity 26.6%; Pred. No. 3.8e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1107
ID ADM05828 standard, protein, 406 AA.
DE Human protein of the invention SEQ ID NO:4513.
PN EP1347046-AA.
                                                                                                                                                                 RESULT 1104
ID ADB64882 standard; protein; 317 AA.
DE Human protein encoded by clone PLACE60070500.
PN EP1308459-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADP55948 standard; protein; 370 AA.
Human PRO protein sequence SEQ ID NO:1924.
WO2004039956-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY08621 standard; protein; 387 AA.
Human secreted protein AJ26_3.
WO9926972-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-SEP-2003.
(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                     (REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-JUL-2001.
(PROT-) PROTEGENE INC.
(SAGA) SAGAMI CHEM RES CENT.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PACE 29-APR-2004.
(FIVE-) FIVE PRIME THERAPEUTICS INC.
(FIVE-) FIVE PRIME THERAPEUTICS INC.
15.1%; Score 288.5; DB 8; Length 244;
ery Match
15.1%; Pred. No. 7.8e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN.
(HRO-) ZH HIROSHIMAZEN SANGYO GIJUTSU SHINKO KI.
(HTRO-) ZH HIROSHIMAZEN SANGYO GIJUTSU SHINKO KI.
15.1%; Score 288; DB 4; Length 192;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; DB 8; Length 402; 4.7e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 15.4%; Score 294; DB 8; Length 344; Local Similarity 29.3%; Pred. No. 4.5e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15.3%; Score 291; DB 7; Length 423; 24.4%; Pred. No. 1e-14;
                                                                                                                                                   Score 295; DB 7; Length 387;
Pred. No. 4.4e-15;
                                                                                                                                                                                                                                                                                                                            Length 387;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 15.5%; Score 295; DB 8; Length 407; Best Local Similarity 27.5%; Pred. No. 4.7e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADMNAYBY Standard; protein; 344 AA.

Human diagnostic and therapeutic pprotein SEQ ID NO:5108.
                                                                                                                                                                                                                ADO41531 standard; protein; 387 AA.
Amino acid sequence of a full length human A34 protein.
WO2004037999-A2.
  26.6%; Pred. No. 4.4e-15;
                                                                                                                                                                                                                                                                                                                          15.5%; Score 295; DB 8; 26.6%; Pred. No. 4.4e-15;
                                                                                                                                                                                                                                                                                                                                                                                       AD041533 standard; protein; 402 AA.
Amino acid sequence of a human A34 protein.
WO2004037999-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         OG-MAY-2004.

(LUDW-) LUDWIG INST CANCER RES.
Query Match 15.5%; Score 295;
BEST Local Similarity 26.6%; Pred. No. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AABB5732 standard; protein; 192 AA.
Rat coxsakie-adenovirus receptor (CAR).
JP2001149095-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADP29419 standard, protein; 244 AA.
Human secreted protein SEQ ID #186.
WO2004035732-A2.
                                      ADC38775 standard; protein; 387 AA.
Human secreted protein #62.
US2002193567-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADQ65260 standard; protein; 407 AA.
Novel human protein sequence #233.
EP1440981-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADC42841 standard; protein; 423 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC78359 standard; protein; 389 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-JUL-2004.
(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                 06-MAY-2004.
(LUDW-) LUDWIG INST CANCER RES.
                                                                                                                       (GEMY) GENETICS INST INC.
ry Match
t Local Similarity 26.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-APR-2003.
(INCY-) INCYTE GENOMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human PRO246 protein.
WO200015796-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2004.
(INCY-) INCYTE CORP.
Best Local Similarity
                                                                                                                                                              Best Local Similarity
RESULT 1113
                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity RESULT 1120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REMAP protein #1. WO2003027228-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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auwsyyyO standard; protein; 413 AA.
Cancer/angiogenesis/fibrosis-related polypeptide, SEQ ID NO:C36O.
WOZO03042661-A2.
                                                                                                                                                                                                                                                                                                                                                                                      ADN39989 standard; protein; 631 AA.
Cancer/angiogenesis/fibrosis-related polypeptide, SEQ ID NO:C359.
WO2003042661-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 377;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 2; Length 318;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-DEC-2000.
(MILL-) MILLERNIUM PHARM INC.
Hery Match 14.2%; Score 271; DB 4; Length 341;
ery Match eimilarity 29.3%; Pred. No. 3.2e-13;
                                                                                                                                                 Length 413;
                                                                                                                                                                                                                                                                                                                              Length 413;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 631;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 631;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 370;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 370;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human secreted protein related protein SEQ ID NO: 136. WO200078808-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AD043537 standard; protein; 370 AA.
Amino acid sequence of a human A33-like 3 protein.
WO2004037999-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 8;
                                                                                                                                               14.9%; Score 285; DB 7; 25.7%; Pred. No. 3.1e-14;
                                                                                                                                                                                                                                                                                                                       14.9%; Score 285; DB 8; 25.7%; Pred. No. 3.1e-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PA (EOSB-) EOS BIOTECHNOLOGY INC.
Query Match 14.9%; Score 285; DB 7;
Best Local Similarity. 25.7%; Pred. No. 5.5e-14;
RESULT 1124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 14.9%; Score 285; DB 8;
Best Local Similarity 25.7%; Pred. No. 5.5e-14;
28.2%; Pred. No. 1.8e-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14.4%; Score 275; DB 4; 29.6%; Pred. No. 1.7e-13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14.3%; Score 272.5; DB 2 28.8%; Pred. No. 2.2e-13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FIVE-) FIVE PRIME THERAPEUTICS INC.
(FIVE-) FIVE PRIME THERAPEUTICS INC.
lery Match
14.9%; Score 284.5; DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MC-NAY-2004.
(LUDW-) LUDWIG INST CANCER RES.
(LUDW-) LUDWIG INST A14%; Score 274; DB 8, lery Match 29.6%; Pred. No. 2e-13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABE6833 standard; protein; 341 AA.
Murine mature INTERCEPT 258 SEQ ID NO: 29.
W02000078808-A1.
                                                                                                                                                                                                              ADR46674 standard; protein; 413 AA.
Cancer-associated protein, SBQ ID 87.
W02004073657-A2.
02-SEP-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADR46673 standard; protein; 631 AA. ADR46673 standard; protein, SEQ ID 86. WO2004073657-A2. (PROT-) PROTEIN DESIGN LABS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human secreted protein; 377 AA. WO2004035732-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAB65905 standard; protein; 370 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW14158 standard; protein; 318 AA
                                                                                                                                                                                                                                                                                                     (PROT-) PROTEIN DESIGN LABS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-MAR-1997.
(LUDW-) LUDWIG INST CANCER RES.
                                                                                                                            (EOSB-) EOS BIOTECHNOLOGY INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                 Best Local Similarity RESULT 1122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 1126
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
RESULT 1123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity RESULT 1129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
RESULT 1128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse A33 antigen.
W09708189-A1.
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AABO8940 standard; protein; 245 AA.
Human secreted protein sequence encoded by gene 13 SEQ ID NO:97.
WO200017222-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY96294 standard; protein; 310 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PD 30-JAN-2003.
PA (ELIL) LILLY & CO ELI.
Query Match
Best Local Similarity 26.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 1144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity RESULT 1145
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RESULT 1147
                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
RESULT 1141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity RESULT 1143
                                                             30-MAR-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-JAN-200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ouery Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 242.5; DB 8; Length 273;
Pred. No. 4.7e-11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 13.0%; Score 247.5; DB 8; Length 228; Best Local Similarity 29.5%; Pred. No. 1.4e-11;
                                                                                                                                                                          AAB65904 standard; protein; 370 AA.

Human secreted protein related protein SEQ ID NO: 134.

BAC200078808-A1.

28-DEC-2000.

(MILL-) MILENNIUM PHARM INC.

ery Match

st Local Similarity 29.3%; Pred. No. 3.6e-13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 294;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-AR-2003.
(HUMA.) HUMAN GENOME SCI INC.
sry Match 13.9%; Score 265; DB 7; Length 294;
it Local Similarity 32.2%; Pred. No. 7.9e-13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 235; DB 8; Length 205;
Pred. No. 1.3e-10;
                                                                                                                     Length 370;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 370;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADO41518 standard; protein; 228 AA.
Amino acid sequence of a human A33-like 3 protein fragment
W02004037999-A2.
                                                                                                                                                                                                                                                                                                RESULT 1132

ID AAB65907 standard; protein; 370 AA.

DE 'Human secreted protein related protein SEQ ID NO: 140.

PD W0200078808-A1.

PD 28-DEC-20007

MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AABG5906 standard; protein; 370 AA.
Human secreted protein related protein SEQ ID NO: 138
WO200078808-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADN02700 standard; protein; 205 AA.
Human receptor and membrane -associated protein #3.
WO2004029218-A2.
                                                                      28-DEC-2000.
(MILL-) MILLENNIUM PHARM INC.
(MILL-) MILLENNIUM PHARM INC.
14.2%; Score 271; DB 4;
cery Match
14.2%; Pred. No. 3.6e-13;
                                                                                                                                                                                                                                                                                                                                                                                          28-DEC-2000.
(MILL-) MILLENNIUM PHARM INC.
(MILL-) MILLENNIUM PHARM INC.
14.2%; Score 270; DB 4;
ery Match
14.2%; Pred. No. 4.3e-13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOZOVAUG-2001.

02-AUG-2001.

(HUMA-) HUMAN GENOME SCI INC.

(HOMA-) HUMAN GENOME SCI INC.

13.9%; Score 265; DB 4;

13.2%; Pred. No. 7.9e-13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14.1%; Score 269; DB 4; 29.0%; Pred. No. 5.2e-13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAU18038 standard; protein; 294 AA.
Human immunoglobulin polypeptide SEQ ID No 183
WO200155315-A2.
                      AAB65832 standard; protein; 370 AA.
Human INTERCEPT 258 SEQ ID NO: 28.
WO200078808-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1135
ID ADB31662 standard; protein; 294 AA.
DE Human novel protein SEQ ID NO 183.
PN US2003077606-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADM87310 standard; protein; 273 AA. Human protein SEQ ID NO:403. WO2004009834-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-MAY-2004.
(LUDW-) LUDWIG INST CANCER RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12.3%;
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                                                                                                                                                                                                                                                                                                                                                                        J-A1.

2000.

LD-) MILLENNIUM

LY MAtch

Best Local Similarity

RESULT 1133

ID AAB65906 stand*

DE Human secr*

PN WOLOON*

PD 28*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-APR-2004.
(INCY-) INCYTE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-JAN-2004.
(NUVE-) NUVELO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
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Best Local Similarity
RESULT 1139
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RESULT 1130
ID AAB65832
DE . Human IN
PN WO200076
PD 28-DEC-.
PA (MILL-)
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AAB00926 standard; protein; 246 AA.
Human secreted protein sequence encoded by gene 13 SEQ ID NO:83.
WO200017222-A1.
                                                                                                                                                            30-MAR-2000.
(HUMA-) HUMAN GENOME SCI INC.
ery Match imilarity 12.0%; Score 228.5; DB 3; Length 246;
(HUMA-) HUMAN GENOME SCI INC.
ry Match 12.0%; Score 228.5; DB 3; Length 245;
t Local Similarity 29.3%; Pred. No. 5.4e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            8; Length 238;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 198.5; DB 6; Length 310;
Pred. No. 2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 4; Length 162;
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                                                                                                                                                                                                                                                                                                                                                                                                   Length 242;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 249;
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Best Local Similarity 31.0%; Pred. No. 7e-10;
RESULT 1142
ID ADR20065 standard; protein; 346 AA.
DE Human immune response associated protein (IRAP), seq:
PP W02004048550-A2.
PD 10-UNN-2004.
PA (INCY-) INCYTE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ë
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11.9%; Score 226.5; DB 8; 24.0%; Pred. No. 1.3e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB65843 standard; protein; 249 AA.
Murine INTERCEPT 258 extracellular domain SEQ ID W0200078808-A1.
                                                                                                                                                                                                                                                                                                                                              18-MAR-2004.
(CYTO-) CYTOS BIOTECHNOLOGY AG.
(CYTO-) CYTOS BIOTECHNOLOGY AG.
11.9%; Score 227; DB 8;
(ery Match 11.9%; Score 227; DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 uvery Match 10.9%; Score 207.5; DB 4
Best Local Similarity 34.8%; Pred. No. 1.5e-08;
RESULT 1146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CYTO-) CYTOS BIOTECHNOLOGY AG.
ry Match
11.8%; Score 225.5; DB 8
t Local Similarity 30.4%; Pred. No. 9.1e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HUMA-) HUMAN GENOME SCI INC.
ry Match
t Local Similarity 29.5%; Pred. No. 7.2e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10.6%; Score 202; DB 4; 27.1%; Pred. No. 7.6e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human junctional adhesion molecule 3 (huJAM3).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABP41477 standard; protein; 226 AA.
Human ovarian antigen HOFNF27, SEQ ID NO:2609
WO200200677-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADL67142 standard; protein; 238 AA.
Mouse B7-HS(ECD) protein SEQ ID NO:12.
WO2004022594-A2.
                                                                                                                                                                                                                                                                               ADL67138 standard; protein; 242 AA.
Human B7-H5(ECD) protein SEQ ID NO:8.
WO2004022594-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAB61417 standard; protein; 162 AA
Human TANGO 244 protein.
WO200100672-A1.
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Length 299;

Length 299;

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AABS3086 standard; protein; 299 AA.
Human angiogenesis-associated protein PRO301, SEQ ID NO:119.
WO200053753-A2.
                              PD 08-JUN-2000.

PA (GFTH) GENENTECH INC.

Query Match 10.3%; Score 196; DB 3; Length 299;

Best Local Similarity 24.8%; Pred. No. 3e-07;

RESULT 1158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10.3%; Score 196; DB 4; Length 299; 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4; Length 299;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAB31202 standard, protein, 299 AA.
Amino acid sequence of human polypeptide PRO301.
WO200077037-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10.3%; Score 196; DB 4; 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                           Score 196; DB 3;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10.3%; Score 196; DB 4; 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10.3%; Score 196; DB 4; 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DETAILS 105-101.
(95-SEP-2001.
(HELL-) HELIX RES INST.
10.3%; Score 196; DB 4;
lery Match
10.3%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10.3%; Score 196; DB 4 24.8%; Pred. No. 3e-07;
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Human immune response protein PRO301 (UNQ264)
WO200119991-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 196; DB 4
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Skin cell protein, SEQ ID NO: 331.
WC20006984-A2.
                                                                                                                                                                                                                                                                                                                                                                                  RESULT 1159
ID AAB80232 standard; protein; 299 AA.
DE Human PRO301 protein.
PN WO200104311-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GENE-) GENESIS RES & DEV CORP LTD.
                                                                                                                                                                                protein; 299 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAM93577 standard; protein; 299 AA.
Human polypeptide, SEQ ID NO: 3365.
EP1130094-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           protein; 299 AA.
SEQ ID NO: 189.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GENE-) GENESIS RES & DEV CORP LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human PRO301 polypeptide sequence. W0200140466-A2.
                                                                                                                                                                         AAY95344 standard; protein; 299
Human PRO301 antitumour protein.
WO200037638-A2.
                                                                                                                                                                                                                                                                ) 29-JUN-2000.

1 (GETH ) GENENTECH INC.

Query Match 10.3%;

Best Local Similarity 24.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PD 18-JAN-2001.

PA (GETH) GENENTECH INC.

Query Match 10.3%;

Best Local Similarity 24.8%;

RESULT 1160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENENTECH INC.
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RESULT 1164
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RESULT 1162
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Best Local Similarity
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RESULT 1163
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB55950 standard;
      WO200032221-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ID AAU00823 8t2
DE Human immuns
PN WO200119991
PD 22-MAR-2001
PA (GETH ) GENI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-DEC-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GETH )
                                                                                                                                                                                                  ADP56683 standard; protein; 310 AA.
Human junction adhesion molecule 3 (huJAM3) full-length protein.
WO2004053058-A2.
                                                                                                                     Score 197.5; DB 3; Length 310;
Pred. No. 2.4e-07;
                                                                                                                                                                                                                                                                                                                                                          Score 197.5; DB 8; Length 310; Pred. No. 2.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PD 04-NOV-1999.

PA (GENE-) GENESIS RES & DEV CORP LTD.

Query Match

Best Local Similarity 24.0%; Pred. No. 3e-07;

RESULT 1156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 299;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         / Match 10.3%; Score 196; DB 2; Length 299; Local Similarity 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10.3%; Score 196; DB 2; Length 299; 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 2; Length 299;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 299;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 196; DB 3; Length 299;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   . (GETH) GENENTECH INC.
Query Match
10.3%; Score 197; DB 2;
Best Local Similarity 24.8%; Pred. No. 2.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10.3%; Score 196; DB 3; 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY23321 standard; protein; 299 AA.
Amino acid sequence of the PRO301 polypeptide.
WO9927098-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 196; DB 2
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human A33 receptor homologue, SEQ ID NO:331.
WO9955865-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY76011 standard; protein; 299 AA.
Human A33 receptor homologue, SEQ ID NO:189.
WO9955865-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB24405 standard; protein; 299 AA.
Human PRO301 protein sequence SEQ ID NO:90.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY13364 standard; protein; 299 AA.
Amino acid sequence of protein PRO301.
WO9914328-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                          AAY08071 standard; protein; 299 AA.
Human PRO307 protein.
WO9914241-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GENE-) GENESIS RES & DEV CORP LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein; 299 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY70670 standard; protein; 299 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SMIK ) SMITHKLINE BEECHAM CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW74464 standard; protein; 29
F11 antigen protein sequence.
WO9902561-A1.
Human IGFAM-6 immunoglobulin. WO200029583-A2.
                                        MOSTON STATE PHARM INC. (INCY-) INCYTE PHARM INC. (INCY-) INCY-) I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-1999.
(GETH) GENENTECH INC.
10.3%;
it Local Similarity 24.8%;
                                                                                                                                                                                                                                                                                               24-JUN-2004.
(RILL) LILLY & CO BLI.
10-BY Match 10.4%;
Best Local Similarity 26.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10.3%;
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GETH ) GENENTECH INC.
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WO200015797-A2.
                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REȘULT 1155
ID · AAY76076 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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Length 299;

Length 299;

Length 299;

Length 299;

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Length 299;
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                                         Length 299;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 299;
                                                                                                                                                                                                                                                                                                                                                                          ABB72150 standard; protein; 299 AA.
Human protein isolated from skin cells SEQ ID NO: 189.
WO200190357-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 24.8%; Pred. No. 3e-07;
RESULT 1181
ID ABO17798 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003032156-A1.
                                                                                                                                                                                                                               ABB72215 standard; protein; 299 AA.
Human protein isolated from skin cells SEQ ID NO: 331
WO200190357-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABB95449 standard; protein; 299 AA.
Human angiogenesis related protein PRO301 SEQ ID NO:
WO200208284-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10.3%; Score 196; DB 6; 24.8%; Pred. No. 3e-07;
                                        10.3%; Score 196; DB 5; 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                  DB 5;
                                                                                                                                                                                                                                                                                                                         10.3%; Score 196; DB 5; 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                              29-NOV-2001.
(GENE-) GENESIS RES & DEV CORP LTD.
10.3*; Score 196; DB 5;
EL Local Similarity 24.8*; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 196; DB 5;
Pred. No. 3e-07;
                                                                                                                                                            (HUMA-) HUMAN GENOME SCI INC.
ry Match 10.3%; Score 196; DB 5
t Local Similarity 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                    (GENE-) GENESIS RES & DEV CORP LTD.
                                                                                          ABG64552 standard; protein; 299 AA.
Human albumin fusion protein #1227.
W0200177137-A1.
18-0CT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABU71610 standard, protein, 299 AA.
Human PRO polypeptide #21.
US2002146709-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABU71465 standard; protein; 299 AA
                       (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10.3%;
24.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human PRO polypeptide #21.

W US2002192659-A1.

19-DEC-2002.

A (GETH ) GENENTECH INC.

Query Match 10.3*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (WARS/) MARSTERS S A.
(PANJ/) PAN J J.
(PADN/) PADNI N F.
(STEP/) STEPHAN J F.
(WAIL/) WILLIAMS P M.
(WOOD/) WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GODOWSKI P J.
GURNEY A L.
HILLAN K J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-OCT-2002.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GERBER H.
GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-FEB-2003.
(GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity RESULT 1182
                                                                                                                                                                                               Best Local Similarity
RESULT 1177
                                                                                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 1178
                                                    Best Local Similarity
RESULT 1176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BAKER K P.
FERRARA N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 1180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GODDARD A.
                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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                                         Query Match
                                                                                                                                                                                    Query Match
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(FERR/)
(GERB/)
(GERR/)
(GODD/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (HILL/)
(MARS/)
(PANJ/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GURN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  лавојаву standard; protein; 299 дд.
Human gene 23 encoded secreted protein fragment, SEQ ID NO:148.
MC200136440-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 лавија втапата, protein; 299 AA.
Human gene 23 encoded secreted protein HACAA29, SEQ ID NO:116.
WO200136440-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human gene 23 encoded secreted protein HACAA29, SEQ ID NO: 86, WO200136440-A1.
                                                      Score 196; DB 4; Length 299;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                         Score 196; DB 4; Length 299;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HUMA-) HUMAN GENOME SCI INC.
10.3%; Score 196; DB 4; Length 299;
it Local Similarity 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 299;
                                                                                                                                                                                                  4; Length 299;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 196; DB 4; Length 299;
Pred. No. 3e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WCACY-2001.
25-MAY-2001.
(HUMA-) HUMAN GENOME SCI INC.
(HTMA-) HUMAN GENOME SCI INC.
10.3%; Score 196; DB 4;
ery Match
10.3%; Score 196; DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-WAY-2001.
(HUNA-) HUMAN GENOME SCI INC.
(ery Match 10.3%; Score 196; DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 196; DB 5;
Pred. No. 3e-07;
                                                                                                                                                                                                  Score 196; DB 4
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10.3%; Score 196; DB 5 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABB84843 standard; protein; 299 AA.
Human PRO301 protein sequence SEQ ID NO:54.
WO200200690-AZ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human polypeptide SEQ ID NO 2666. WO200190304-A2. 29-MOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                 Human novel protein; 299 AA. WO200155437-A2.
                                                                                                         AAU14405 standard; protein; 299 AA.
Human novel protein #276.
WO200155437-A2.
                                                                                                                                                                                                                                              AAU14404 standard; protein; 299 AA.
Human novel protein #275.
WO200155437-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABG64551 standard; protein; 299 AA.
Human albumin fusion protein #1226.
WOZ00177137-A1.
18-OCT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein; 299 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                10.3%;
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                                PA (GETH ) GENENTECH INC.
Query Match 10.3%;
Best Local Similarity 24.8%;
RESULT 1167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENENTECH INC.
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.uA-) HUMAN GENON.
.ry Match
Best Local Similarity .RESULT 1173
ID ABB90290 stander
PN WA20010-
PN WA20010-
PN WA20010-
PN WA20010-
PN PD 29-
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                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
RESULT 1169
ID AAU14168 standard,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
RESULT 1171
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RESULT 1172
                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAE03840 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                              02-AUG-2001.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                      02-AUG-2001.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HYSE-) HYSEQ INC.
PD 14-SEP-2000.
PA (GRTT)
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(GETH ) GEN
                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                  Query Match
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Length 299;

DB 6; Length 299;

RESULT 1183
ID ABO2517
DE Novel h
PN US20030
PD 27-FEB-

RESULT 1185

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Score 196; DB 6; Length 299;
Pred. No. 3e-07;
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Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABU67160 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301 US2003032062-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABUG7028 standard; protein; 299 AA.
Human secreted/transmembrane, PRO, protein SEQ ID 366.
US2003032155-A1.
                                                                                                                                                                                                                                                                                                              ABU72059 standard, protein, 299 AA.
Novel human secreted and transmembrane protein PRO301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human secreted / transmembrane polypeptide PRO301
                      ABO25023 standard; protein; 299 AA.
Human secreted/transmembrane protein (PRO) #183.
US2003036179-A1.
                                                                                                                                                                                                                                                                                                                                                                                                        10.3%; Score 196; DB 6; 24.8%; Pred. No. 3e-07;
                                                                                                          Score 196; DB 6;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ouery Match 10.3%; Score 196; DB 6; Best Local Similarity 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-OCT-2002.
(GETH ) GENENTECH INC.
2ry Match 10.3%; Score 196; DB 6;
2ry Match 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 196; DB 6
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human junctional adhesion molecule 1 (huJAM1) WO2003008541-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 196; DB 6
Pred. No. 3e-07;
                                                                                                                    Lest Local Similarity 24.8%; Pred. No. 3e-RESULT 1193

ID ABG64519 standard; protein; 299 AA.

EBS Human secreted/transmembrane protein, #23.

PN US2002160374-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABU67365 standard; protein; 299 AA.
Human secreted protein PRO301.
US2003023054-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human A-33 related antigen PRO301. US2002182206-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            299 AA
                                                   10.3%;
24.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 10.3%;
Best Local Similarity 24.8%;
RESULT 1201
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ELIL ) LILLY & CO ELI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-FEB-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 1197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 1199
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                                                                                                                                                                                                                                                                                                                                                  US2002177165-A1.
28-NOV-2002.
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   RESULT 1192
                                                                                                                                                                                                                                                                                                                                                                                                                           Length 299;
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Pred. No. 3e-07;
                                                                                                                           10.3%; Score 196; DB 6; Length 299; 24.8%; Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
                                  ABO25173 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.
27-PEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABO01794 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2002197671-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel human secreted and transmembrane protein PRO301 US2003032063-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABO47382 standard; protein; 299 AA.
Human secreted/transmembrane polypeptide PRO301
US2003044839-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABUS9833 standard; protein; 299 AA.
Novel secreted and transmembrane protein PRO301
US2003017563-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                             DB 6;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                             ABU71911 standard; protein; 299 AA.
Human secreted/transmembrane protein PRO301.
US2003003530-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABUS4367 standard; protein; 299 AA.
Human secreted/transmembrane protein PRO301.
US2000132240-A1.
19-SEP-2002.
(GETH ) GENENTECH INC.
                                                                                                                                                                             ABUB1052 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003004311-A1.
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23-JAN-2003.
(GETH ) GENENTECH INC.
10.3%;
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1 (GETH) GENENTECH INC.

QUERY MATCH 10.3%;

BEST LOCAL SIMILARITY 24.8%;
                                                                                                                                                                                                                                                                                                                                                                                                        (GETH ) GENENTECH INC.
ry Match 10.3%;
t Local Similarity 24.8%;
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Best Local Similarity 24.8%;
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Best Local Similarity 24.8%;
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Local Similarity 24.8%;
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Human PRO polypeptide #183.
US2003036180-A1.
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Best Local Similarity 24.8%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                          (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                         (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
 Best Local Similarity
                                                                                                                                         Best Local Similarity RESULT 1184
                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
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Best Local Similarity
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                                                                                                                               Query Match
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Length 299;

Length 299;

Length 299

Length 299;

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Length 299;

DB 6;

RESULT 1190

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Human PRO polypeptide #183.
US2003082704-A1.
01-MAY-2003.
                                                                                               Score 196; DB 6; Length 299;
Pred. No. 3e-07;
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Pred. No. 3e-07;
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(GETH) GENENTECH INC.
ery Match 10.3%; Score 196; DB 6; Length 299;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 6; Length 299;
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Pred. No. 3e-07;
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                                                                                                                                                                                                                                                     6; Length 299;
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Pred. No. 3e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     JT 1204
JT 1204
JT 1204
Novel human secreted and transmembrane protein PRO301.052003022328-A1.
30-AN-2003.
(GETH ) GENENTECH INC.
10.3%; Score 196; DB 6; Lengt etcy Match
11.3%; Pred. No. 3e-07;
ABU69642 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003017463-A1.
                                                                                                                                                                                                                                                                                                                             Human secreted / transmembrane polypeptide PRO301.
US2003027143-A1.
                                                                                                                                                                                                                                                     Score 196; DB 6,
Pred. No. 3e-07;
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(GETH) GENENTECH INC.
10.3%; Score 196; DB 6,
tr Local Similarity 24.8%; Pred. No. 3e-07;
                                                                                                                                                   ABU79802 standard; protein; 299 AA.
Human secreted/transmembrane protein PRO301.
US2003032057-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADB29324 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003092002-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADA76316 standard, protein, 299 AA.
Human PRO polypeptide #183.
US2003073212-A1.
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ID ADA61589 standard; protein; 299 AA.
DE , Homo sapiens.
PN US2003049816-A1.
                                                                                                                                                                                                                                                                                                         ABO14824 Btandard; protein; 299 AA
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13-FBE-2003.
(GETH ) GENENTECH INC.
10.3%;
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06-PEB-2003.
(GETH ) GENENTECH INC.
10.3%;
                                                 PD 23-JAN-2003.
PA (GETH ) GENENTECH INC.
Query March
Best Local Similarity 24.8%;
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(GETH ) GENENTECH INC.
bry Match
10.3%;
st Local Similarity 24.8%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
RESULT 1204
ID AD45885 standard; px
DE Novel human secreted
PN US2003022328-A1.
PD 30-0AN-2003.
PA (GETH ) GENENTECH INC
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Best Local Similarity
RESULT 1203
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Best Local Similarity
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Length 299;
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                   Length 299
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Best Local Similarity 24.8%; Pred. No. 3e-07;
RESULT 1218
ID ADAS5842 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003082693-A1.
                                                                       ADA86394 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US200308211-A1.
01-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                PD 17-APR-2003.
PA (GETH ) GENENTECH INC.
Query March 10.3%; Score 196; DB 6;
Best Local Similarity 24.8%; Pred. No. 3e-07;
                 10.3%; Score 196; DB 6; 24.8%; Pred. No. 3e-07;
                                                                                                                                                                       DB 6;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
                                                                                                                                                                       Score 196; DB 6
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 24.8%; Score 196; DB RESULT 1215
DD ABO32776 standard; protein; 299 AA.
DE Human secreted/transmembrane protein PRO301.
PN US2003045693-A1.
DD 06-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADA18180 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003039971-A1.
                                                                                                                                                         Best Local Similarity 24.8%; Pred. RESULT 1212

ID ADB15958 standard; protein; 299 AA. DE Human PRO polypeptide #183.

PN US2003087350-A1.
                                                                                                                                                                                                                                                                                                                                                                                ADA47744 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003073215-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADA67539 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003068795-Al.
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Human PRO polypeptide #183.
US2003068794-A1.
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Human PRO polypeptide #183.
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24.8%;
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24.8%;
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Query March 10.3%;
Best Local Similarity 24.8%;
RESULT 1213
                                                                                                                                                     (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Query Match
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(GETH ) GENENTECH INC.
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                                 Best_Local Similarity RESULT 1211
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Best Local Similarity
RESULT 1217
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RESULT 1219
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(GETH ) GENENTECH
                                                                                                                                                                                                                                                                                       08-MAY-2003
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                   Query Match
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01-MAY-2003
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                                                   Score 196; DB 6; Length 299;
Pred. No. 3e-07;
                                                                                                                                                                                                 Query Match 10.3%; Score 196; DB 6; Length 299; Best Local Similarity 24.8%; Pred. No. 3e-07;
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Pred. No. 3e-07;
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                                                                                                                                                                                                                                                ADAB1797 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.
US2003087345-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADA91791 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.
US2003082694-A1.
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Novel human secreted and transmembrane protein PRO301
US2003073211-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 196; DB 6;
Pred. No. 3e-07;
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Human secreted/transmembrane polypeptide PRO301
US2003044844-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 196; DB.6;
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 196; DB 6
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADA16155 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003049621-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 10.3%; Score 196; Local Similarity 24.8%; Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                       Human PRO polypeptide #183.
US2003087349-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABO346346 standard; protein; 299 AA.
Human PRO polypeptide #21.
US2003044793-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADB14854 standard, protein, 299 AA.
Human PRO polypeptide #183.
US2003087351-A1.
                                                                                                        protein; 299 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                          USZUCZ-C
08-MAY-2003.
(GETH ) GENENTECH INC.
10.3%;
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                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 24.8%;
RESULT 1222
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Best Local Similarity 24.8%;
                                                   Query Match
Best Local Similarity 24.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 24.8%;
                                                                                                                         Human PRO polypeptide #183
US2003082763-A1.
                 01-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                           01-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                         08-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 1225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                      ADA79358 standard;
US2003082705-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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Score 196; DB 6; Length 299;
Pred. No. 3e-07;
                           Length 299;
                                                                                                                                                Score 196; DB 6; Length 299;
Pred. No. 3e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10.3%; Score 196; DB 6; Length 299; 24.8%; Pred. No. 3e-07;
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                                                                                                                                                                                Novel human secreted and transmembrane protein PRO301 US2003082691-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                            ABO43331 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301 US2003044945-A1.
                                                                                                                                                                                                                                                                        Score 196; DB 6;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 196; DB 6;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                 Score 196; DB 6
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
                           Score 196; DB (
Pred. No. 3e-07,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADA42300 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003054401-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human PRO polypeptide SEQ ID NO 366. 192500307713-A1.
                                                                                                                                                                                                                                                                                                           Human PRO polypeptide #183. US2003082710-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human PRO polypeptide #183. US2003082701-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JA74492 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003068798-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA75212 standard; protein; 299 AA.
Human PRO polypeptide #183.
17-2003073216-A1.
17-APR-2003.
                                                                    ADA94030 standard; protein; 299 AA
Human PRO polypeptide #183.
US2003077722-Al.
                                                                                                                                                                                                                           Chery Match 10.3%;
Best Local Similarity 24.8%;
RESULT 1229
                                                                                                                                                  10.3%;
                                                                                                                                                                                                                                                                                                                                                        PD 01-MAY-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 24.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10.3%;
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PA (GETH ) GENENTECH INC.
Query Match 10.3%;
Best Local Similarity 24.8%;
RESULT 1233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 10.3%;
Best Local Similarity 24.8%;
RESULT 1236
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                                                                                                                     24-APR-2003. (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                             Best Local Similarity
RESULT 1230
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Length 299;

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Length 299;
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Novel human secreted and transmembrane protein PRO301
US2003082765-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10.3%; Score 196; DB 6; 24.8%; Pred. No. 3e-07;
                                                                                                                                  10.3%; Score 196; DB 6; 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                      DB 6;
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Best Local Similarity 24.8%; Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10.3%; Score 196; DB 6; 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                     10.3%; Score 196; DB 6 24.8%; Pred. No. 3e-07;
                                        Human PRO polypeptide #183.
US2003092147-A1.
                                                                                                                                                                       Human PRO polypeptide #183.
US2003096386-Al.
AMAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB34173 standard; protein; 299 AA. Human PRO polypeptide SEQ ID NO 366. US2003077714-A1.
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US2003082760-Al.
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US2003082702-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADA96502 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003082690-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADA95950 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003082759-Al.
                                                                                                                                                                                                                                                                                                                     ADA61026 standard; protein; 299 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10.3%;
24.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PD 01-MAY-2003.

PA (GETH) GENENTECH INC.

Query Match 10.3%;

Best Local Similarity 24.8%;

RESULT 1253
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Best Local Similarity 24.8%;
                                                                                                                                                                                                                                                                                      24.8%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                        13-MAR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GETH ) GENENTECH INC.
          Best Local Similarity
RESULT 1247
ID ADB26811 standard; pr.
DE Human PRO polypeptide
PN US2003092147-A1.
PD 15-MAY-2003.
PA (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                       (GETH ) GENENTECH INC.
                                                                                                                 (GETH ) GENENTECH INC
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                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
RESULT 1250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
                                                                                                                                            Best Local Similarity RESULT 1248
                                                                                                                                                                                                                                                                                 Best Local Similarity
RESULT 1249
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RESULT 1255
                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
US2003049817-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-APR-2003
                                                                                                                                    Query Match
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 Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-AFK-2005.
(GETH ) GENENTECH INC.
iry Match
ir Toral Similarity 24.8%; Pred. No. 3e-07;
            Score 196; DB 6; Length 299;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                    Score 196; DB 6; Length 299;
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
                                                                                                                                                   Length 299,
                                                              ADA85290 standard; protein; 299 AA.

Novel human secreted and transmembrane protein PRO301
US2003082695-A1.
                                                                                                                                                                                                 ADA04738 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003082708-A1.
                                                                                                                                                   DB 6;
                                                                                                               01-MAY-2003.
(GETH ) GENENTECH INC.
iry Match 10.3%; Score 196; DB 6;
iry Match 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADB25285 standard; protein; 299 AA.
Human PRO polypeptide SEQ ID NO 366.
US2003077115-A1.
24-APR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADA75764 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003082703-A1.
                                                                                                                                                                                                                                                                                                                                ABO17514 standard; protein; 299 AA.
Human PRO polypeptide #21.
US2003064367-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human PRO polypeptide #183.
17-APD-2002
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Human PRO polypeptide #183.
US2003073210-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADA93461 standard, protein, 299 AA.
Human PRO polypeptide #183.
US2003077721-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein; 299 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US2005000
01-MAY-2003.
(GETH ) GENENTECH INC.
MATCh ... 1arity 24.8%;
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01-MAY-2003.

(GETH ) GENENTECH INC.

10.3$;
               10.3%;
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RESULT 1241
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ry Match
t Local Similarity 24.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADA80522 standard; protein;
Human PRO polypeptide #183
US2003082761-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                      03-APR-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                     (GETH ) GENENTECH INC.
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PA . (GETH ) GENENTECH INC.
                                                                                                                                                              Best Local Similarity RESULT 1239
                           Best Local Similarity RESULT 1238
                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity RESULT 1245
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                   Query Match
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ADA42726 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003054351-A1.
            RESULT 1265
                                                                                                                            DB 7; Length 299;
                                                                                                                                                                                                                                                               Score 196; DB 7; Length 299;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 196; DB 7; Length 299;
Pred. No. 3e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADA46437 standard; protein; 299 AA.

Novel human secreted and transmembrane protein PRO301.

N US2003054916-A1.

O 20-MAR-2003.

A (GETH ) GENENTECH INC.

Ouery Match

Dest Local Similarity 24.8%; Pred. No. 3e-07;
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(GETH ) GENENTECH INC.

ery Match

10.3%; Score 196; DB 7; Length 299;
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Novel human secreted and transmembrane protein PRO301.
US2003082709-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 196; DB 7;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 7;
Best Local Similarity 24.8%; Pred. No. 3e-07; RESULT 1256
                                                                                                                            Score 196; DB 7
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                  Score 196; DB 7
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 196; DB 7
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADA41876 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003082540-A1.
GGTH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADAIG579 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003039969-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human secreted/transmembrane protein, #25. Was 032003017498-A1.
                                        ADA77523 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003068797-A1.
                                                                                                                                                                           ADB18263 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003077710-A1.
                                                                                                                          10.3%;
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Best Local Similarity 24.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   V Match 10.3%;
Local Similarity 24.8%;
                                                                                                                                                                                                                                                                                                                                                                 01-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-FEB-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                          (GETH ) GENENTECH INC.
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                                                                                                                                      Best Local Similarity RESULT 1257
                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 1260
ID ADA13008 standard;
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Best Local Similarity
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                                                                                                                                                                                                                              24-APR-2003
                                                                                            10-APR-2003
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DB 7; Length 299;
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Pred. No. 3e-07;
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Novel human secreted and transmembrane protein PRO301.
US203087344-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADA88601 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003073213-A1.
                                                                                                                                                                                                                                                                                                                                               Score 196; DB 7;
Pred. No. 3e-07;
                                                                                   Score 196; DB 7;
Pred. No. 3e-07;
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Best Local Similarity 24.8%; Pred. No. 3e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 7;
                                                                                                                                                                     DSzvo...
01-MAY - 2003.
(GETH ) GENENTECH INC.
nerv Match 10.3%; Score 196; DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 196; DB 7
Pred. No. 3e-07;
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Pred. No. 3e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ID ABO19860 standard; protein; 299 AA.
DB Human secreted/transmembrane protein PRO302
PN . US203044902-A1.
PD 06-MAR-2003.
D 20-WAR-2003

PA (GETH ) GENENTECH INC.

Query Match

Best Local Similarity 24.8%; Pred. No.

RESULT 1266

ID ADB28467 standard; protein; 299 AA.

DE Human PRO polypeptide #183.

PN US2003082699-A1.

PA (FEW.
                                                                                                                                                                                                                                                                                                                                                                                 Human PRO polypeptide #183.
US2003059909-A1.
                                                                                                                                                                                                                                                             ADB29019 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003082706-A1.
01-MAY-2003.
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US2003022239-Al.
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Human PRO polypeptide #21.
                                                                                                                                                                                                                                                                                            US2003002...
01-MAY-2003.
(GETH ) GENENTECH INC.
10.3%; Sr
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24.8%;
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 1269
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 1273
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RESULT 1267
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RESULT 1268
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Length 299;

Length 299;

Length 299;

DB 7;

Length 299;

Length 299;

Length 299;

DB 7;

Length 299;

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10.3%; Score 196; DB 7; Length 299; 24.8%; Pred. No. 3e-07;
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Novel human secreted and transmembrane protein PRO301.
US2003082696-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADB47114 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003082687-A1.
                                                                                                                                                                                                                                                                                                                                                             Novel human secreted and transmembrane protein PRO301
US200308254-A1.
                                        10.3%; Score 196; DB 7; 24.8%; Pred. No. 3e-07;
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                                                                                                                                                                                   Score 196; DB 7;
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Best Local Similarity 24.8%; Fred. No. 3e-07;
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Best Local Similarity 24.8%; Pred. No. 3e-07;
RESULT 1289
                                                                                                                                                                                                                                     ADB77645 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003077654-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human secreted/transmembrane protein, #25. US2003082542-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human PRO polypeptide SEQ ID NO 366. 24-app
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human PRO polypeptide #183.
US2003082697-A1.
                                                                                        ADB90390 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003082762-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADB74781 standard; protein; 299 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10.3%;
                                                                                                                                                                                   10.3%;
24.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 10.3%;
Best Local Similarity 24.8%;
RESULT 1288
                                                           24.88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
     01-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
                                                     Best Local Similarity
RESULT 1284
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
RESULT 1290
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RESULT 1285
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Best Local Similarity
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(GETH ) GEN
                                                                                                                                                01-MAY-2003
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                                          Query Match
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Beet Local Similarity 24.8%; Pred. No. 3e-07;
RESULT 1281
                                                                                                                                                                                                Score 196; DB 7; Length 299;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                            10.3%; Score 196; DB 7; Length 299; 24.8%; Pred. No. 3e-07;
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(GETH ) GENENTECH INC.
rry Match
rr Tocal Similarity 24.8%; Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
                                                           Score 196; DB 7; Length 299;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADA92143 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.
US2003082712-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADB30658 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.
US2003082766-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADB38106 standard; protein; 299 AA.

Novel human secreted and transmembrane protein PRO301.
US2003087347-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADB66578 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.
US2003082689-A1.
                                                                                                                                                                                                                                                                                                                                                                                     Human PRO polypeptide SEQ ID NO 366 24-APB-2702
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 24.8%; Score RESULT 1283
ID ADB89658 standard; protein; 299 AA. DE Human PRO polypeptide #183.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -DBB15406 standard; protein; 299 AA. Human PRO polypeptide #183. US2003087352-A1.
                                                                                                              protein; 299 AA.
                                                                                                                                                                                                                                                   ADB22848 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003077711-A1.
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                                                           10.3%;
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Best Local Similarity 24.8%;
RESULT 1279
                                                                                                                                                                             PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 24.8%;
RESULT 1276
                                                                                                                                 Human PRO polypeptide #183
US2003068793-A1.
                                                                                                                                                                                                                                                                                                           24-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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GETH ) GENENTECH INC.
Query Match 1
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PD 08-MAX-2003.
PA (GETH ) GENENTECH INC.
                                          (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
23-A1.
2003.
...H ) GENENTECH .
..ry Match
Best Local Similarity
RESULT 1275
ID ADA66987 stand**
DE Human PRO r
PN US2003 r
PD 10**
                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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Best Local Similarity
RESULT 1278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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Length 299;

RESULT 1293

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(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
10.3%; Score 196; DB 7; Length 299;
10.3%; Score 196; DB 7; Length 299; 24.8%; Pred. No. 3e-07;
                                                                                                                                                         10.3%; Score 196; DB 7; Length 299; 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                   Length 299;
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Novel human secreted and transmembrane protein PRO301.
US2003092106-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AUC/1954 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.
US203092107-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC59933 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003092105-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 10.3%; Score 196; DB 7; Best Local Similarity 24.8%; Pred. No. 3e-07; RESULT 1309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10.3%; Score 196; DB 7; 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                     DB 7;
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15-MAY GENENTECH INC.
(GETH ) GENENTECH INC.
Herv Match 10.3%; Score 196; DB 7;
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PA (GETH ) GENENTECH INC.

QUELY MATCH

10.3%; Score 196; DB 7, Best Local Similarity 24.8%; Pred. No. 3e-07; RESULT 1306
                                                                                                                                                                                                                                                                                                 Query Match 10.3%; Score 196; DB 7 lbest Local Similarity 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 196; DB 7
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human secreted/transmembrane protein, #25.
US2003054441-A1.
                                                                                                                                                                                                                  ADC20851 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003049677-A1.
                                                                                                                                                                                                                                                                                                                                                                             ADC40736 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003054400-A1.
                                                        ADC29320 standard; protein; 299 AA.
Muman secreted(transmembrane protein, #25.
US2003049676-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC33841 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003073077-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human secreted/transmembrane protein, #25. US2003073079-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADC12911 standard; protein; 299 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PD 17-APR-2003.

PA (GETH ) GENENTECH INC.

QUERY MATCh

10.3%;

Best Local Similarity 24.8%;

RESULT 1308
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                         (GETH ) GENENTECH INC.
                Best Local Similarity RESULT 1302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
                                                                                                                                                         Query Match
Best Local Similarity
RESULT 1303
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Best Local Similarity
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ID ADC19393 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
    Query Match
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                    10.3%; Score 196; DB 7; Length 299; 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                DB 7; Length 299;
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1 (GETH ) GENENTECH INC.

Query Match 10.3%; Score 196; DB 7; Length 299;

Best Local Similarity 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                          Length 299;
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Pred. No. 3e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel human secreted and transmembrane protein PRO301
US2003082692-A1.
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Pred. No. 3e-07;
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                                                                                                                                                                              Score 196; DB 7
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADC39627 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003059828-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 1300
ID ADC18969 standard, protein, 299 AA.
DE Human secreted/transmembrane protein, #25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human secreted/transmembrane protein, #25.
US2003059772-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADC40141 standard; protein; 299 AA. Human secreted/transmembrane protein, #25. US2003059829-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC34265 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003036094-Al.
                                                                                                                                                                                                                                                                                                                                                                                        Human PRO polypeptide SEQ ID NO 366.
US2003077720-A1.
                                                                            ADB33931 standard; protein; 299 AA. Human PRO polypeptide SEQ ID NO 366 US2003077716-A1.
                                                                                                                                                                                                                                ADB35035 standard; protein; 299 AA.
Human PRO polypeptide SEQ ID NO 366
US2003077718-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                299 AA
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DE Human secreted/transmembrane pro US2003069829-A1.

PD 27-MAR-2003.

PA (GETH ) GENENTECH INC.

Query Match

10.34; Sc
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Local Similarity 24.8%;
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Best Local Similarity 24.8%;
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Best Local Similarity 24.8%;
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1 (GETH ) GENENTECH INC.

10-31;

Best Local Similarity 24.81;
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                 24-APR-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                           (GETH ) GENENTECH INC.
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  (GETH ) GENENTECH INC
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                      Query Match
Best Local Similarity
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20-FEB-2003.
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ADC58517 standard; protein; 299 AA.
Novel human secreted and transmembrane protein Seg ID366.
US2003087346-A1.
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     RESULT 1320
                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 196; DB 7; Length 299;
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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                                                                                                                                    Length 299
                                                                                                                                Best Local Similarity 24.8%; Pred. No. 3e-07; REGIT 1312

ID ADC57294 standard; protein; 299 AA.

DE Novel human secreted and transmembrane protein Seq ID366.

PN US200308736-A1.

PD 08-MAY-2003.

PA (GETH ) GENENTECH INC.
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Novel human secreted and transmembrane protein Seq ID366.
US2003007364-A1.
(8-NAX-2003.
(GETH ) GENENTECH INC.
                                ADC52940 standard; protein; 299 AA.
Novel human secreted and transmembrane protein Seg ID366.
US2003087365-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADC59069 standard; protein; 299 AA.
Novel human secreted and transmembrane protein Seg ID366.
US2003087359-A1.
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Novel human secreted and transmembrane protein Seg ID366.
US2003087360-A1.
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Novel Muman secreted and transmembrane protein PRO301
082003087367-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADC50960 standard; protein; 299 AA.

Novel human secreted and transmembrane protein PRO301
US2003087361-A1.
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
Best Local Similarity 24.8%; Fred. No. 3e-07; RESULT 1311
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Human PRO polypeptide #183.
US2003087362-A1.
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08-MAY-2003.
(GETH ) GENENTECH INC.
watch 10.3%; St.
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08-MAY-2003.

(GETH ) GENENTECH INC.

MATCh ''larity 24.8%;
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(GETH) GENENTECH INC.

10.3%;

Query Match 10.3%;
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38-MXY-2003.
(GETH ) GENENTECH INC.
10.3%;
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(GETH ) GENENTECH INC.
10.3%;
FY MATCH Similarity 24.8%;
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A (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 24.8%;
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(GETH) GENENTECH INC.
Query Match
Best Local Similarity 24.8%;
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RESULT 1314
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Length 299;
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Pred. No. 3e-07;
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Novel human secreted and transmembrane protein PRO301
US2003087354-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADC80551 standard; protein; 299 AA.

Novel human secreted and transmembrane protein PRO301
US2003092103-A1.
                                                                                                                                                                                                                          ADD03191 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 24.8%; Score 196; DB 7; RESULT 1326
DE Human DDA ---
                                                 Score 196; DB 7;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 7;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
                                                                                                                                                                             DB 7;
                                                                                                                                                                                                                                                                                                       10.3%; Score 196; DB 7; 24.8%; Pred. No. 3e-07;
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Best Local Similarity 24.8%; Pred. No. 3e-07;
RESULT 1322
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Pred. No. 3e-07;
                                                                                          ADCI2363 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003082541-A1.
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15-00m 200 Human PRO polypeptide #183.
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Human PRO polypeptide #183.
US2003194773-A1.
US2003ve...
08-MAY-2003.
(GETH ) GENENTECH INC.
March 10.3%; Sr
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24.8%;
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24.8%;
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Best Local Similarity 24.8%;
RESULT 1328
                                                                                                                                                                                                                                                       US2003092104-A1.
15-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                              01-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 1329
                                                           Best Local Similarity
RESULT 1321
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Best Local Similarity
RESULT 1325
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RESULT 1323
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RESULT 1324
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Score 196; DB 7; Length 299;
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Local Similarity 24.8%; Pred. No. 3e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 299,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADC79999 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.
US2003087358-A1.
                                                                                                                                               ADD1343 standard; protein; 299 AA.
Human secreted/transmembrane PRO polypeptide #27
US2003105011-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADD11303 standard; protein; 299 AA.
Human secreted/transmembrane PRO polypeptide #27
US2003105013-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 196; DB 7;
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
10.3%; SCORE 196; DB 7;
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Pred. No. 3e-07;
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US2003104469-Al.
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ID ADD03924 standard, protein, 299 AA.

DE Human secreted/transmembrane protein, #25.

PN US2003104381-A1.
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16-0CT-2002
                                                                                                                                                                                                                                                                                       protein; 299 AA.
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Human PRO polypeptide #183.
US2003194775-A1.
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2003.
) GENENTECH INC.
10.3%; S.
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OS-JUN-2003.
(GETH ) GENENTECH INC.
MATCh '17rity 24.8%;
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                                                              16-OCT-2003.
(GETH ) GENENTECH INC.
2ry Match 10.3%;
3t Local Similarity 24.8%;
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Local Similarity 24.8%;
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                                                                                                                                                                                                                                                                                ACC77939 standard; protein.
Human PRO polypeptide #183
US2003194771-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GETH ) GENENTECH INC.
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30-OCT-2003.
(GETH ) GENENTECH INC.
10.3%; Score 196; DB 7; Length 299;
ery Match 10.3%; Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Novel human secreted and transmembrane protein PRO301. US2003203438-A1.
                                                                                                                                                                                                                                                                                                                                                                                                  ADD53612 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.
US2003203437-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADD54183 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003203432-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADD37096 standard; protein; 299 AA.
Human secreted/transmembrane PRO polypeptide #27
US2003105012-A1.
                                                                                                                                                                                                                                                                                                                                               Score 196; DB 7;
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
                                                                      Score 196; DB 7
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Human PRO polypeptide #183.
US2003194792-A1.
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10c2003203430-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADD02567 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003203431-A1.
                                                                                                                      protein; 299 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein; 299 AA
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Human PRO polypeptide #183.
                 US2003202.
30-CCT-2003.
(GETH ) GENENTECH INC.
10.3%; SC
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30-OCT-2003.
(GETH ) GENENTECH INC.
10.3%; Sr
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24.8%;
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Query Match 10.3%;
Best Local Similarity 24.8%;
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Query March
Best Local Similarity 24.88;
RESULT 1340
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Best Local Similarity 24.8%;
                                                                                                                 ADD52320 standard; protein,
Human PRO polypeptide #183
US2003194769-A1.
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Human PRO polypeptide #183
US2003194779-A1.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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Best Local Similarity
RESULT 1339
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RESULT 1343
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10.3%;
24.8%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                       (GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 1361
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                                                Best Local Similarity
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RESULT 1363
       16-OCT-2003
                                                                                                                                                                                                                                                06-NOV-2003
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                                                                   RESULT 1357
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                                                   Score 196; DB 7; Length 299;
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Novel human secreted and transmembrane protein PRO301.
US2003194767-A1.
                                                                                                                                                                                                                                                                                                                                      ADE32307 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003194765-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 7;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Best Local Similarity 24.8%; Pred. No. 3e-07;
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US2003199023-A1.
                                                                                             ADD91396 standard, protein; 299 AA.
Human PRO polypeptide #183.
US2003199055-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE22239 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003199056-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADD79463 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003203428-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADE41999 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003194772-A1.
16-OCT-2003.
(GETH ) GENENTECH INC.
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1D ADD91948 standard; protein; 299 AA.
DE Human PRC polypeptide #183.
PN US2003199053-A1.
                                                                                                                                                                                                                   ADE04010 standard; protein; 299 AA. Human PRO polypeptide #183. US2003199057-A1.
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...dard; protein,
...d199055-A1.
23-OCT-2003.
PA (GETH) GENENTECH INC.
Ouery Match
Best Local Similarity 24.8%; Pre
RESULT 1349
ID ADE04010 standard; protein
PR UNS2003199057
PD 23-OCT
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16-007-2003.
(GETH ) GENENTECH INC.
MATCh ... 134;
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23-007-2003.
(GETH ) GENENTECH INC.
10.3%;
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23-OCT-2003.
(GETH ) GENENTECH INC.
MATCh 1-11arity 24.8%;
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(GETH ) GENENTECH INC.
(ATT ) GENENTECH INC.
(ATCh ) 1943;
                                                                                                                                                                                                                                                                                              10.3%;
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RESULT 1354
                                                                 24.8%;
                                                     10.3%;
                       23-OCT-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
.30-Al.
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.ry Match
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Best Local Similarity .RESULT 1348
ID ADD91396 stand>
DE Human PRO r
PN US20031
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Best Local Similarity
RESULT 1352
ID ADD79463 standard, pl
DE Human PRO polypeptide
PN US200320348-A1.
PD 30-CCT-2003.
PA (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                       Best Local Similarity RESULT 1350
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Best Local Similarity
RESULT 1351
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Length 299;
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 Length 299;
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                                                  ADE33963 standard, protein, 299 AA.

Novel human secreted and transmembrane protein PRO301
US2003194791-A1.
                                                                                                                                                                                                                                                      Best Local Similarity 24.8%; Score 196; DB 7; RESULT 1359

TESULT 1359

DE Human PRO polypeptide #183.

PN US2003194768-A1.

PA GERTY.
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Best Local Similarity 24.8%; Pred. No. 3e-07;
RESULT 1362
ID ADE18920 standard; protein; 299 AA.
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10.3%; Score 196; DB 7; 24.8%; Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Human secreted/transmembrane protein, #25.
US2003077583-A1.
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US2003199026-Al.
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US2003199025-Al.
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RESULT 1365
ID ADE22791 standard; protein; 299 AA.
BN US2003199064-A1.
PD 23-OCT-2003.
                                                                                                                                      Query Match
Best Local Similarity 24.8%; Pred.
RESULT 1358
ID ADD80015 standard; protein; 299 AA.
DE Human PRO Polypeptide #183.
PN US2003207417-A1.
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Human PRO polypeptide #183.
US2003199033-A1.
23-OCT-2003.
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Length 299;

Query Match

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Score 196; DB 7; Length 299;
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
                                                ADG21157 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003207384-A1.
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Novel'human secreted and transmembrane protein PRO301
US2003207381-A1.
                                                                                                                                     Score 196; DB 7;
Pred. No. 3e-07;
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Human secreted/transmembrane polypeptide PRO301
US2003170721-A1.
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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US2003039972-Al.
                                                                                                                                                                                    ADF97492 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003207370-A1.
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   10.3%;
24.8%;
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PA (GETH ) GENENTECH INC.
QUEYY MATCh
BEST LOCAL Similarity 24.8%;
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24.8%;
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Query Match
Best Local Similarity 24.8%;
RESULT 1383
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Best Local Similarity 24.8%;
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Best Local Similarity 24.8%;
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(GETH ) GENENTECH INC.
                                                                                                  06-NOV-2003.
(GETH ) GENENTECH INC.
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Human PRO301 protein.
US2003171568-A1.
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(GURN/) GURNEY A L.
(NAPL) NAPIER M A.
(TUMA/) TUMAS D.
(WOOD/) WOOD W I.
              Best Local Similarity
RESULT 1375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1-SEP-2003.
(ASHK/) ASHKENAZI A.
(FONG/) FONG S.
                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 1378
                                                                                                                                                      Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
RESULT 1379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                 Length 299;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Best Local Similarity 24.8%; Score 196; DB 7; Length 299;
RESULT 1372
                                                                                                                                                      Length 299
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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                                                                                                                                                                                                    ADE32859 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003194766-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG21516 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.
US2003207355-A1.
                 Score 196; DB 7;
Pred. No. 3e-07;
                                                                                                                                                      DB 7;
                                                                                                                                                    Score 196; DB 7
Pred. No. 3e-07;
                                                            ADD78909 standard, protein, 299 AA. Human PRO polypeptide #183. US2003203429-A1. 30-OCT-2003. (GFH.) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human PRO polypeptide #183.
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182003194777-Al.
                                                                                                                                                                                                                                                                                                                                        ADE42551 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003199032-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADD89595 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003199028-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADE04678 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003199034-A1.
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                10.3%;
                                                                                                                                                  10.3%;
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Best Local Similarity 24.8%;
RESULT 1368
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Best Local Similarity 24.8%;
RESULT 1369
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1 (GETH) GENENTECH INC.

100-X Match

Best Local Similarity 24.8%;
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A (GETH) GENENTECH INC.

10-USY MATCH

Best Local Similarity 24.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE40879 standard; protein;
Human PRO polypeptide #183.
US2003199031-A1.
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                                                                                                                                                                                                                                                    16-OCT-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                            23-OCT-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
GENENTECH INC
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                                                                                                                                                                Best Local Similarity RESULT 1367
              Query Match
Best Local Similarity
RESULT 1366
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DB 7; Length 299;

Length 299;

Length 299;

Length 299;

Length 299;

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Best Local Similarity
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Best Local Similarity
RESULT 1398
                                                                              Best Local Similarity RESULT 1393
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RESULT 1401
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RESULT 1397
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                                                                      Query Match
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Best Local Similarity 24.8%; Pred. No. 3e-07;
RESULT 1388
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Pred. No. 3e-07;
                                                                                   Score 196; DB 7; Length 299;
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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                                                                                                                                                                                                                                           RESULT 1385
ID AD164067 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.
PN US2003207385-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1387

ID ADI63515 standard; protein; 299 AA.

DE Novel human secreted and transmembrane protein PRO301.

PN US2003207387-A1.
ADH55848 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.
US2001207379-A1.
                                                                                                                                                                                                                                                                                                                                                                                      AD165016 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003207386-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADH81929 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003207388-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADH81377 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003207377-A1.
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Human secreted/transmembrane protein, #25.
US2003054349-A1.
                                                                                                                   Human secreted/transmembrane protein, #25. US200305452-A1.
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06-NOV-2003.
(GETH ) GENENTECH INC.
10.3%; S'
                                                                                                                                                                                                                                                                                               US200320.
06-NOV-2003.
(GETH ) GENENTECH INC.
10.3%;
                                      06-NOV-20us.
(GETH ) GENENTECH INC.
10.3%;
Pry Match 10.3%;
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RESULT 1392
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Best Local Similarity 24.8%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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Best Local Similarity
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Best Local Similarity
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Length 299;
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Novel human secreted and transmembrane protein PRO301.
US2003087353-A1.
08-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                        ADNI5393 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.
US2003087356-A1.
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Novel human secreted and transmembrane protein PRO301.
US2003087357-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADC81103 standard; protein; 299 AA.

Novel human secreted and transmembrane protein PRO301.
US2003092115-A1.
                                                                                                                                   ADNI6574 standard, protein; 299 AA.
Novel human secreted and transmembrane protein PRO301 US2003087385-A1.
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                                                                                                                                                                                                                                10.3%; Score 196; DB 7; 24.8%; Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                       Score 196; DB 7;
Pred. No. 3e-07;
                                                                             10.3%; Score 196; DB 7; 24.8%; Pred. No. 3e-07;
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Human secreted/transmembrane protein, #25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADE79197 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US20031335025-A1.
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Human PRO polypeptide #183.
US2003100087-A1.
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30-0CT-2003.
(GETH ) GENENTECH INC.
10.3%;
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Query Match 10.3%;
Best Local Similarity 24.8%;
RESULT 1395
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Human PRO polypeptide #183.
US200320340-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                  (GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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05-JUN-2003
                                                                                                                                                                                                       Score 196; DB 8; Length 299; Pred. No. 3e-07;
                                                       10.3%; Score 196; DB 8; Length 299; 24.8%; Pred. No. 3e-07;
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Pred. No. 3e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 8; Length 299
                                                                                                                                                                                                                                                                                                                                                                                                          ADE41304 standard; protein; 299 AA.
Human secreted/transmembrane PRO polypeptide #27
US2003100497-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADE41186 standard; protein; 299 AA.
Human secreted/transmembrane polypeptide PRO301
US2003104558-A1.
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(GETH) GENENTECH INC.

Query Match

10.3%; Score 196; DB 8

Best Local Similarity 24.8%; Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
                                                                                                                                                                                                                                                         ADE73297 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003129592-A1.
                                                                                                     ADE75767 standard; protein; 299 AA. Human PRO polypeptide #183. US2003211571-A1. 13-NOV-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 1406

ID, ADE23895 standard; protein; 299 AA.

DE Human PRO polypeptide #183.

PN US2003092110-A1.

PD 15-MAY-2003.

PA (GETH) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JAE23343 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003092108-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADE24518 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003092111-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADD87363 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003203439-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADE89229 standard, protein, 299 AA. Human PRO polypeptide #183. US2003199062-A1. CS3-OCT-2003. (GETH ) GENENTECH INC.
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                                                                                                                                                                                                      Query Match 10.3%;
Best Local Similarity 24.8%;
RESULT 1403
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Best Local Similarity 24.8%;
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Best Local Similarity 24.8%;
                                                                         24.8%;
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1 (GETH ) GENENTECH INC.

QUEYY MATCH 10.3%;

BEST LOCAL Similarity 24.8%;
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(GETH ) GENENTECH INC.
US2003130489-A1.
10-JUL-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-OCT-2003.
(GETH ) GENENTECH INC.
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                                                                     Best Local Similarity
RESULT 1402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 1407
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                                                       Query Match
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RESULT 1404
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PD 23-OCT-2003.

PA (GETH ) GENENTECH INC.

Query Match 10.3%; Score 196; DB 8; Length 299;

Best Local Similarity 24.8%; Pred. No. 3e-07;

RESULT 1416
                                                                                                                                                                   uery Match 10.3%; Score 196; DB 8; Length 299;
Best Local Similarity 24.8%; Pred. No. 3e-07;
SULT 1412
                       Score 196; DB 8; Length 299;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                      Score 196; DB 8; Length 299;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 8; Length 299;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10.3%; Score 196; DB 8; Length 299; 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 196; DB 8; Length 299;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 299;
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Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 196; DB 8
Pred. No. 3e-07;
                                                               ADE99386 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003211576-A1.
                                                                                                                                                                                                                                              ADE18368 standard, protein; 299 AA. Human PRO polypeptide #183. US2003194794-A1. 16-OCT-2003. (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADE95249 standard, protein; 299 AA.
Human PRO polypeptide #183.
US2003199052-A1.
                                                                                                                                                                                                                                                                                                                                                                                                               ADE88677 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003199054-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADE94697 standard; protein; 299 AA. Human PRO polypeptide #183. US2003199027-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE91108 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003199061-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADE93359 standard, protein; 299 AA.
Human PRO polypeptide #183.
US2003199060-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADF34940 standard, protein; 299 AA.
Human PRO polypeptide #183.
US2003199029-Al.
23-0CT-2003.
Query Match 10.3%;
Best Local Similarity 24.8%;
                                                                                                                                                                                                                                                                                                                                                    Query Match 10.3%;
Best Local Similarity 24.8%;
RESULT 1413
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PA (GETH ) GENENTECH INC.

Query March

Best Local Similarity 24.8%;

RESULT 1414
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PA (GETH ) GENENTECH INC.

QUETY MATCh 10.3%;

Best Local Similarity 24.8%;

RESULT 1415
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Best Local Similarity 24.8%;
                                                                                                                                            07-AUG-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-OCT-2003.
(GETH ) GENENTECH INC.
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(GETH) GENENTECH INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity RESULT 1417
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RESULT 1428
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(GETH ) GENENTECH INC.

(ery Match 10.3%; Score 196; DB 8; Length 299;

(ery Match 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 vuery Match
Beet Local Similarity 24.8%; Pred. No. 3e-07;
RESULT 1424
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(GETH) GENENTECH INC.
ery Match
-- '---- Similarity 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PD 25-SEP-2003.

PA (GETH) GENENTECH INC.

Query Match 10.3*; Score 196; DB 8; Length 299;

Best Local Similarity 24.8*; Pred. No. 3e-07;
                   Score 196; DB 8; Length 299;
Pred. No. 3e-07;
                                                                                                                                                                        DB 8; Length 299;
                                                                                                                                                                                                                                                                                                                          Length 299;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE91703 standard; protein; 299 AA.

Novel human secreted and transmembrane protein PRO301.
US2003199058-A1.
23-OCT-2003.
(GETH.) GENENTECH INC.
                                                                                                                                                                                                                             ADE92255 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.
US2003199051-A1.
                                                                                                                                                                                                                                                                                                                          Score 196; DB 8;
Pred. No. 3e-07;
                                                                                                                      US4000---
13-NOV-2003.
(GETH ) GENENTECH INC.
Orv Match 10.3%; Score 196; DB B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 196; DB 8,
Pred. No. 3e-07;
                                                                         ADE98505 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003211569-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADE99932 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003211568-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADG40402 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003225253-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADF73796 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003180312-A1.
                                                                                                                                                                                                                                                                                                                                                                              ADE90556 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003199063-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein; 299 AA
                   10.3%;
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Best Local Similarity 24.8%;
RESULT 1423
                                                                                                                                                                                                                                                                                     23-OCT-2003.
(GETH ) GENENTECH INC.
iry Match 10.3%;
it Local Similarity 24.8%;
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US2003207352-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GODD/) GODDARD A.
(GODD/) GODDWSKI P J.
(GOND/) GODDWSKI P J.
(GURH/) GURNER J P.
(WILL/) WILLIAMS P M.
(WODD/) WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                       23-OCT-2003.
(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
                               Best Local Similarity
RESULT 1420
                                                                                                                                                                                      Best Local Similarity RESULT 1421
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Best Local Similarity
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Best Local Similarity
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Best Local Similarity
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                     Query Match
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Length 299;
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ADG22068 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.
US2003207360-A1.
                                                                                                                                                                                                                                                                                                                                                                                   ADG24261 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003207426-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Luery Match 10.3%; Score 196; DB 8;
Best Local Similarity 24.8%; Pred. No. 3e-07;
RESULT 1434
                                                                              Score 196; DB 8;
Pred. No. 3e-07;
                                                                                                                                                                                                             DB 8;
                                                                                                                                                                                                                                                                                                                                          DB 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10.3%; Score 196; DB 8; 24.8%; Pred. No. 3e-07;
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Pred. No. 3e-07;
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QUETY MATCH
10.3%; Score 196; DB 8
Best Local Similarity 24.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                      10.3%; Score 196; DB 8 24.8%; Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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US2003207359-A1.
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US2003207375-A1.
                                                                                                                              ADG20138 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003207376-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human PRO polypeptide #183.
US2003207353-A1.
                                                                                                                                                                                                                                                            protein; 299 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADF98615 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003208055-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADG03446 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003207351-A1.
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                                              PD 06-NOV-2003.
PA (GETH ) GENENTECH INC.
QUery Match 10.3%;
Best Local Similarity 24.8%;
RESULT 1429
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PA (GETH) GENENTECH INC.

Query Match 10.3%;

Best Local Similarity 24.8%;

RESULT 1436
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 1433
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Best Local Similarity
RESULT 1437
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RESULT 1431
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Query
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                                                                                  Score 196; DB 8; Length 299;
Pred. No. 3e-07;
                                                                                                                                                                                                                     Score 196; DB 8; Length 299;
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                              ADG08372 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003207424-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel human secreted and transmembrane protein PRO301.
US203207389-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 196; DB 8;
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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                                                                                                                                ABF73372 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003166051-A1.
                                                                                                                                                                                                                                                                                                                                         (GETH ) GENENTECH INC.
10.3%; Score 196;
it Local Similarity 24.8%; Pred. No. 3
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US2003207371-Al.
ADG19478 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003207425-Al.
                                                                                                                                                                                                                                                                     ADG13315 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003207357-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADG15542 standard, protein, 299 AA.
Human PRO polypeptide #183.
US2003219885-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADG06125 standard, protein, 299 AA.
Human PRO polypeptide #183.
US2003207374-A1.
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US2003207423-Al.
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04-SEP-2003.
(GETH ) GENENTECH INC.
Match 10.3%;
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Local Similarity 24.8%;
                                                                                    Query Match 10.3%;
Best Local Similarity 24.8%;
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Local Similarity 24.8%;
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Local Similarity 24.8%;
                                                   06-NOV-2003.
(GETH ) GENENTECH INC.
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                                                                                                                                                                                                                                   Best Local Similarity RESULT 1439
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Length 299;
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Novel human secreted and transmembrane protein PRO301.
US2003207427-A1.
                                                                                                                            ADG07196 standard, protein; 299 AA.

Novel human secreted and transmembrane protein PRO301
US2003207350-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                            ADG55243 standard, protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003194778-A1.
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Novel human secreted and transmembrane protein PRO301 US2003207428-A1.
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Novel human secreted and transmembrane protein PRO301
                                                                                                                                                                                                                                                                ADG07748 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003207356-A1.
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Novel human secreted and transmembrane protein PRO301
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US2002207362-A1.
                                                                        Score 196; DB 8;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                        Score 196; DB 8;
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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PA (GETH ) GENENTECH INC.

Query Match 10.3*; Score 196; DB 8

Best Local Similarity 24.8*; Pred. No. 3e-07;

RESULT 1448
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Pred. No. 3e-07;
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Human secreted/transmembrane protein, #25.
US2003027145-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10.3%; Score 196;
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               US2003207
06-NOV-2003.
(GETH ) GENENTECH INC.
10.3%; S'
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Ouery Match
Best Local Similarity 24.8%;
RESULT 1449
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24.8%;
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Human PRO polypeptide #183.
US2003207358-A1.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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Best Local Similarity
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                                                                         Query Match
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Length 299;

Length 299;

Length 299;

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10.3%; Score 196; DB 8; Length 299; 24.8%; Pred. No. 3e-07;
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Best Local Similarity 24.8%; Score 196; DB 8; Lengt
RESULT 1468
ID ADG54139 standard; protein; 299 AA.
DE Novel human secreted and transmembrane protein PRO301.PD NG5003207416-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADG56347 standard, protein, 299 AA.
Novel human secreted and transmembrane protein PRO301.
US2003207366-A1.
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Novel human secreted and transmembrane protein PRO301.
US2003207429-A1.
                                                                                                                                                                                                         ADH11989 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003207419-A1.
                                                                                                                                                                                                                                                                                                                                         ADG52411 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003207414-A1.
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ADH12613 standard; protein; 299 AA.
US2003207378-A1.
                                                                 ADG63621 standard; protein; 299 AA.
Human secreted/transmembrane polypeptide PRO301
US2003180796-A1.
25-SEP-2003.
                                                                                                                                                        10.3%; Score 196; DB 8; 24.8%; Pred. No. 3e-07;
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Pred. No. 3e-07;
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192003194793-A1.
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US200302231-Al.
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16-OCT-2003.
(GETH ) GENENTECH INC.
10.3%; S
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 1470
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Best Local Similarity
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                                                                                                                                                                      Local Similarity
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RESULT 1473
                             Best Local Similarity RESULT 1465
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                  Query Match
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Pred. No. 3e-07;
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Pred. No. 3e-07;
                                                   Score 196; DB 8; Length 299;
Pred. No. 3e-07;
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Pred. No. 3e-07;
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Novel human secreted and transmembrane protein PRO301.
US2003207415-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AUG/1473 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.
US200207421-A1.
                                                                                                                                                                                                                                          ADGS8555 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301.052003207368-A1.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                        Novel human secreted and transmembrane protein PRO301.
US2003207420-A1.
                                                                                             ADG55795 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003207365-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADG58003 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003207363-A1.
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADG81660 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003207805-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADH30622 standard; protein; 299 AA.
Human PRO polypeptide #183.
US2003077723-A1.
24-APR-2003.
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06-NOV-2003.
(GETH ) GENENTECH INC.
"arch "reity 24.8%;
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06-FEB-2003.
(GETH ) GENENTECH INC.
MAtch '12tity 24.8%;
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06-NO-2003.
(GETH ) GENENTECH INC.
MATCH '134;
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(GETH ) GENENTECH INC.
iry Match 10.3%;
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24.8%;
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RESULT 1462
                                GENENTECH INC.
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                                                                 Best Local Similarity RESULT 1456
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Best Local Similarity
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Best Local Similarity
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US2003207364-A1.
                  06-NOV-2003
(GETH ) GEN
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Length 299;

DB 8;

Length 299;

Score 196; DB 8; Pred. No. 3e-07;

Length 299;

Length 299;

Length 299;

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US2003207361-A1.
10.3%; Score 196; DB 8; Length 299; 24.8%; Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Pred. No. 3e-07;
                                                                                                 ADG54691 standard, protein, 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003207367-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                    ADG59731 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301 US2003207369-A1.
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Pred. No. 3e-07;
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Pred. No. 3e-07;
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Human secreted/transmembrane protein, #25.
US2004005553-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADH07286 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2004006211-A1.
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Human secreted/transmembrane protein, #25.
US2003215904-A1:
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US2004005665-A1.
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Human PRO polypeptide #183.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USJUCTORY OF THE TRACE OF THE T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 24.8%;
RESULT 1476
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24.8%;
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Local Similarity 24.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 10.3%;
Best Local Similarity 24.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADH43487 standard; protein
Human PRO polypeptide #27.
US2003224984-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-JAN-2004.
DESN/) DESNOYERS L.
(GODD/) GODDARD A.
(GODO/) GODOWSKI P J.
(GURN/) GURNEY A L.
(WAILL/) WILLIAMS P M.
(WOOD/) WOOD W I.
                                                                                                                                                                                                                                  06-NOV-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-NOV-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-DEC-2003.
(GETH ) GENENTECH INC.
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(DESN) DESNOTERS L.
(GODD/) GODDARD A.
(GODO/) GODOWSKI P J.
(GURN/) GURNEY A L.
(MATH/) MATHER J P.
(WILL/) WILLIAMS P M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
RESULT 1479
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Query Match
Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GURN/)
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GG-NOV-2003.
(GETH) GENENTECH INC.
ery Match 10.3%; Score 196; DB 8; Length 299;
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PD (GETH ) GENENTECH INC.

Query Match

10.3$; Score 196; DB 8; Length

Best Local Similarity 24.8$; Pred. No. 3e-07;

RESULT 1485

ID ADGOS989 standard; protein; 299 AA.

DE Novel human secreted and transmembrane protein PRO301.

PN US2004009548-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADI15369 standard, protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003207382-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADI14701 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003207383-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADG09246 standard; protein; 299 AA.

Novel human secreted and transmembrane protein PRO301
US2004009547-A1.
                                                                                                                                                                                                                 Score 196; DB 8;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                  Score 196; DB 8;
Pred. No. 3e-07;
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Local Similarity 24.8%; Pred. No. 3e-07;
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Best Local Similarity 24.8%; Pred. No. 3e-07;
                                                                                         Human secreted/transmembrane protein, #25. US2003152999-A1.
                                                                                                                                                                   PD 14-AUG-2003.

PA (GETH ) GENENTECH INC.

Query Match

10.34; Score 196; DI

Best Local Similarity 24.84; Pred. No. 3e-18

RESULT 1483

ID AD165321 standard; protein; 299 AA.

DE Human secreted/transmembrane protein, #25.
                                                                                                                                                                                                                                                                                                                                                                                                                                           ADI37584 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003096340-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADH97380 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003190610-AI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADI65748 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003148371-A1.
                                                                                                                                                                                                                                                                                                                                         PD 07-AUG-2003.

PA (GETH ) GENENTECH INC.

Query March 10.3%;

Best Local Similarity 24.8%;

RESULT 1484
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(GETH ) GENENTECH INC.
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RESULT 1486
                                                          Best Local Similarity
RESULT 1482
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RESULT 1487
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                                                  Query Match
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PA (GETH) GENENTECH INC.
Query Match
10.3%; Score 196; DB 8; Length 299;
Best Local Similarity 24.8%; Pred. No. 3e-07;
RESULT 1491
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(GETH) GENENTECH INC.
ery Match 10.3%; Score 196; DB 8; Length 299;
st Local Similarity 24.8%; Pred. No. 3e-07;
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A (GETH ) GENENTECH INC.
Query Match 10.3%; Score 196; DB 8; Length 299;
Best Local Similarity 24.8%; Pred. No. 3e-07;
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(GETH ) GENENTECH INC.
"" Matrh 10.3%; Score 196; DB 8; Length 299;
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                                                                                                                                                                                                                                                                                                                   DB 8; Length 299;
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Novel human secreted and transmembrane protein PRO301.
US2004039164-A1.
                                                                                                                                                                                                                                                                                                                                                                          ADI18296 standard; protein; 299 AA.
Novel human secreted and transmembrane protein PRO301
US2003207349-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 196; DB 8;
Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                   10.3%; Score 196; DB 8 24.8%; Pred. No. 3e-07;
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Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADJ99548 standard, protein, 299 AA.
Human secreted/transmembrane protein, #25.
US2003187238-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADLO8741 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003186358-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADM25082 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2003096233-A1.
                                                                                                 ADH60491 standard; protein; 299 AA.
Human secreted/transmembrane protein, #25.
US2004023331-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-MAR-2004.
(UVNY ) UNIV NEW YORK STATE RES FOUND.
ery Match
10.3%; Score 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADK40844 standard; protein; 299 AA. Human platelet F11 receptor #1. US6699688-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein; 299 AA
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2003:
GENENTECH INC.
10.3%; Sr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human JAM-1 protein sequence. WO2004003145-A2.
                                                                                                                                                                          (GODD/) GODDARD A. (GODD/) GODDARD A. (GODN/) GODDWSKI P J. (GUN/) GURNEY A L. (MATH/) MATHER J P. (WILL/) WILLIAMS P M. (WODD/) WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-OCT-2003.
(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 1498
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Best Local Similarity
RESULT 1492
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Best Local Similarity
RESULT 1496
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Best Local Si
RESULT 1493
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Best Local S
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ID ADLO874
DE Human B
PN US20031
PD 02-OCT-
PA (GETH )
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Best Local Similarity 24.8%; Pred. No. 3e-07;
RESULT 1499

ID ADM29812 standard; protein; 299 AA.
DE Human secreted/transmembrane protein, #25.
PN US2003190611-A1.
PD 09-0CT-2003.
PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 24.8%; Pred. No. 3e-07;
RESULT 1500

ID ADQ5890 standard; protein; 299 AA.
DE T cell activation associated protein #34.
PN WO2004058805-A2.
PN WO2004058805-A2.
PA (ASAH-) ASAHI KASEI PHARMA CORP.
Query Match
ACH (ASAH-) ASAHI KASEI PHARMA CORP.
Rest Local Similarity 24.8%; Pred. No. 3e-07;
Result Activation associated protein #34.
PA (ASAH-) ASAHI KASEI PHARMA CORP.
Rest Local Similarity 24.8%; Pred. No. 3e-07;
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Length 2458;
    Length 2458;
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Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 2458;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACA63509 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACA71673 standard; cDNA; 2458 BP.
Human secreted and transmembrane polypeptide PRO363 cDNA
US2002177553-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel human secreted and transmembrane protein PRO363 US2003003531-A1.
                                                                                                                                                                                                                                                                                         63
                                                                                                                                                                                                                                                              ABL95592 standard; cDNA; 2458 BP.
Human angiogenesis related cDNA PRO363 SEQ ID NO:
WO200208284-A2.
    DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 6;
                                                                                                                                                                                           DB 6;
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80
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    100.0%; Score 2458; 100.0%; Pred. No. 0;
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Pred. No. 0;
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No. 0;
                                                                                                                                                                                           Score 2458;
Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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cDNA encoding human PRO polypeptide #194.
US2003036180-A1.
                                                                     ABL88103 standard; cDNA; 2458 BP.
Human PRO363 cDNA sequence SEQ ID NO:63.
WO200200690-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Score 100.0%; Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACA64513 standard; cDNA; 2458 BP.
                                                                                                                                                                                         100.0%;
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RESULT 9
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A (GETH ) GENENTECH INC.
Query Match 100.0%;
Best Local Similarity 100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (PAGN/) PAGNI N F.
(STBP/) STEPHAN J F.
(WATA/) WATANABE C K.
(WILL/) WILLIANABE P M.
(WOOD/) WOOD W I.
                                                                                                                                                                  (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                   GERBER H.
GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                              GENENTECH INC
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GODOWSKI P J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MARSTERS S A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                 Best Local Similarity RESULT 6
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HILLAN K J.
                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                        FERRARA N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
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US2003017563-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PAN J
                                                                                                                                            03-JAN-2002
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GERB/)
GERR/)
(GODD/)
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(PAON/)
(STEP/)
                                                                                                                                                                                                                                                                                                                                                                                    BAKE/)
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GenCore version 5.1.6
6, 2005, 04:10:11; Search time 1295 Seconds
(Without alignments)
11236.080 Million cell updates/sec
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                                                                                                                                            ....acagagcgggactccgtctc 2458
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                                                                                                                                                                                                                                          8780412
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Human PRO363 protein UNQ318 encoding cDNA SEQ ID NO:86.
WO200053758-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAC78473 standard; cDNA; 2458 BP.
Human PRO363 (UNQ318) nucleotide sequence SEQ ID NO:58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAS21437 standard; cDNa; 2458 BP.
Human cDNA sequence encoding for PRO363 polypeptide.
WO200140466-A2.
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Human PRO363 nucleotide sequence SEQ ID NO:502.
WO200073454-Al.
                                                                                                                                       Sequence: 1 gcgccgggagcccatctgcc.....aca Scoring table: IDENTITY NUC Gapext 1.0 Gapext 1
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(GETH ) GENENTECH INC.
100.0%; Score 2458;
ETY Match 100.0%; Pred. No. 0;
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(GETH ) GENENTECH INC.
iry Match 100.0%; Score 2458;
iry Match 100.0%; Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Human PRO363 nucleotide sequence.
WO9946281-A2.
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(GETH) GENENTECH INC.
Match '100.0%;
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2458
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(GETH ) GENENTECH INC
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Best Local Similarity
RESULT 3
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Best Local Similarity
RESULT 4
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Best Local Si
RESULT 5
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RESULT 13

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DB 8; Length 2458;
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13-FEB-2003.
(GETH) GENENTECH INC.
100.0%; Score 2458;
Query Match
100.0%; Pred. No. 0;
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Best Local Similarity 100.0%; Pred. No. 0;
RESULT 29
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Pred. No. 0;
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US2003073212-A1.
                                                                                                                                                                                   ABX17255 standard; cDNA; 2458 BP. Human PRO polynucleotide #140. US2002123463-A1.
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US2003054517-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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RESULT 28
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Best Local Similarity
RESULT 30
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Human PRO363 cDNA.
US2003050448-A1.
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                                                                                                                                                                                                                                                                                               RESULT 25
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Human cDNA encoding a secreted/transmembrane protein, SEQ ID 387.
US2003032155-Al.
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                                                                                              DB 8; Length 2458;
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Novel human secreted and transmembrane protein PRO363 cDNA, 052003088063-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA US2003022187-Al.
30-JAN-2003.
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Human cDNA encoding secreted/transmembrane protein PRO363
US2003004102-A1.
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Novel human secreted or transmembrane protein PRO5723 DNA
US20031027985-A1.
06-PEB-2003.
                                                                                                                                                                                                                                                                                                                                                                                      Human secreted/transmembrane protein (PRO) cDNA #194.
US2003036179-A1.
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Human sectreted/transmembrane protein cDNA, #176.
US2002142961-A1.
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8
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PN . US2003027162-A1.
PD 06-FEB-2003.
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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 2458;
               ABX92313 standard; CDNA; 2458 BP.
CDNA encoding human PRO363 polypeptide.
US2002169284-Al.
14-NOV-2002.
(GETH) GENENTECH INC.
18-Y Mach 1 100.0%; Score 2458; St. Local Similarity 100.0%; Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
                                                                                                                                                                                                                                                 ACD44481 standard; cDNA; 2458 BP. cDNA encoding human PRO363 polypeptide.
US2002127576-Al.
                                                                                                                                           ABX80972 standard; cDNA; 2458 BP.
                                                                                                                                                                                                                                                                                                                                                                         ACD41988 standard; cDNA; 2458 BP
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12-SEP-2002.
(GETH ) GENENTECH INC.
100.0%;
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03-OCT-2002.
(GETH ) GENENTECH INC.
100.0%;
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20-FEB-2003.
(GETH ) GENENTECH INC.
100.0%;
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08-MAY-2003.
(GETH) GENENTECH INC.
100.0%;
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RESULT 16
ID ACD41988 standard; cl
DE Human secreted/transn
PN US2003036179-A1.
PD 20-FEB-2003.
PA (GETH ) GENENTECH INC
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Best Local Similarity
RESULT 20
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Best Local Similarity
RESULT 15
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RESULT 19

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20-MAR-2003.
(GETH ) GENENTECH INC.
100.0%; Score 2458; DB 9; Length 2458;
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Pred. No. 0;
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                                                                                                                                                                                                                                                                     Length 2458;
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                                                                                          DB 8; Length 2458;
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Novel human secreted and transmembrane protein PRO363 CDNA.
2500302238-A1.
ACA93171 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA US2003011476-A1.
                                                                                                                                                                                                                                                                                                                                   ACA68110 standard, cDNA, 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2002177164-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human secreted and transmembrane polypeptide PRO363 cDNA US2002197615-A1.
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Human PRO polynucleotide #194.
US2003068795-A1.
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                                                                      Query Match
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                                                                                                                     DB 9; Length 2458;
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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 2458; DB 9; Length 2458;
ery Match 100.0%; Pred. No. 0;
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Human cDNA encoding secreted/transmembrane polypeptide PRO363.
US2003054404-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AUA10486 standard; CDNA; 2458 BP.
Human CDNA encoding secreted/transmembrane protein, PRO363.
US2003059831-A1.
                                                                                                                                                                ADB19395 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003068796-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                       ADA86415 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003082711-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADA38013 standard; cDNA; 2458 BP.

Human cDNA encoding secreted/transmembrane protein PRO363
US2003008297-A1.
09-JAN-2003.
(GETH ) GENENTECH INC.
100.0%; Score 2458; DB 9; Length;
t.cocal Similarity 100.0%; Pred. No. 0;
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08-MAY-2003.
(GETH) GENENTECH INC.
(GETH) 100.0%; Score 2458;
ery Match 100.0%; Pred. No. 0;
                                                                               13-MAR-2003.
(GETH ) GENENTECH INC.
iry Match 100.0%; Score 2458;
iry Match 100.0%; Pred. No. 0;
     100.0%; Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
                                                                                                                                                                                                                                                                                                  ADB27936 standard; cDNA; 2458 BP.
cDNA encoding human PRO polypeptide #194.
US2003082704-A1.
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US2003073215-A1.
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Human PRO polynucleotide #194.
US2003087350-A1.
                                  ADA61610 standard; cDNA; 2458 BP.
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10-APR-2003.
(GETH.) GENENTECH INC.
100.0%;
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01-MAY-2003.
(GETH ) GENENTECH INC.
"...ch 100.0%;
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17-APR-2003.
(GETH ) GENENTECH INC.
"Arch "Tetty 100.04;
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Best Local Similarity RESULT 32
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Best Local Similarity
RESULT 39
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                                                   Homo sapiens.
US2003049816-A1.
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Best Local Si
RESULT 35
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Best Local Si
RESULT 36
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ID ADA47
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100.0%; Score 2458; DB 9; Length 2458; 100.0%; Pred. No. 0;
                                      DB 9; Length 2458;
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                                                                                                                                                                                                                                ADA85863 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003082693-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADA87518 standard; cDNA; 2458 BP. Novel human secreted and transmembrane protein PRO363 cDNA. US2003087345-Al.
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Novel human secreted and transmembrane protein PRO363 cDNA.
US2003082694-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADA28138 standard; cDNA; 2458 BP.
Human cDNA encoding secreted/transmembrane protein PRO363
US2003054359-A1.
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(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
100.0%; Bred. No. 0;
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                                     Score 2458;
Pred. No. 0;
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100.0%; Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
                                                                                     ADB10567 standard; cDNA; 2458 BP.

CDNA encoding human PRO polypeptide #194.

US2003068794-A1:
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CDNA encoding human PRO363 polypeptide.
US2003054987-A1.
20-MAR-2003.
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US2003087349-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADA97075 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003082705-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADA79379 standard; cDNA; 2458 BP. Human PRO polynucleotide #194. US2003082763-A1.
                INC.
100.0%;
EV 100.0%;
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RESULT 43
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                     Local Similarity
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Best Local Similarity
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10-APR-2003.
(GETH ) GENENTECH
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(GETH) GENENTECH INC. 100.0%; Score 2458; 2ry Match 100.0%; Pred. No. 0;

Query Match Best Local Similarity RESULT 51

ADB14875 standard; cDNA; 2458 BP. Human PRO polynucleotide #194. US2003087351-A1.

S2003uc.)8-MAY-2003. (GETH) GENENTECH INC. 100.0%;

Best Local Similarity RESULT 52

Query Match

Query Match Best Local Similarity RESULT 53

Score 2458; Pred. No. 0;

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DB 9; Length 2458;
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USAUCAUS.

(GETH) GENENTECH INC.

100.0%; Score 2458;

100.0%; Pred. No. 0;
Score 2458;
Pred. No. 0;
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Best Local Similarity 100.0%; Pred. No. 0;
RESULT 65
                                                                                                                                                                                                                                                                               100.0%; Score 2458; 100.0%; Pred. No. 0;
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Pred. No. 0;
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cDNA encoding human PRO polypeptide #194.
US2003073214-A1.
                                                                                                                                                                                                                                                                                                                                   ADB24746 standard; cDNA; 2458 BP.
Human PRO polynucleotide SEQ ID NO 387.
US2003077713-A1.
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US2003082761-Al.
                                                                                                                                                                                    ADA74513 standard; cDNA; 2458 BP. Human PRO polynucleotide #194. US2003068798-A1.
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US2003073216-A1.
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Human PRO polynucleotide #194.
US2003082701-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADA85311 standard; cDNA; 2458 BP.
                                                        ADA94718 standard; cDNA; 2458 BP
 100.0%;
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A (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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                 Best Local Similarity RESULT 60
                                                                                                                                               Best Local Similarity RESULT 61
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Best Local Similarity
RESULT 67
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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 2458; DB 9; Length 2458;
ery Match 100.0%; Pred. No. 0;
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Human cDNA encoding secreted/transmembrane polypeptide PRO363.
US2003055216-Al.
                     DB 9; Length 2458;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 2458;
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                                                                                                                                                                                                                       ADA24597 standard; cDNA; 2458 BP.

Novel human secreted and transmembrane protein PRO363 cDNA.
US2003050241-A1.
13-MAR-2003.
(GETH ) GENENTECH INC.
100.0%; Score 2458; DB 9; Length 24
st Match
st Local Similarity 100.0%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACD98617 standard, cDNA, 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003044945-Al.
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Novel human secreted and transmembrane protein PRO363 cDNA.
US2003025040-A1.
13-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                            ADB18836 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003073211-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADB19947 standard; CDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 CDNA, 012003082691-A1.
                                                                                                                                                                      DB 9;
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100.0%; Score 2458; 100.0%; Pred. No. 0;

ADA94051 standard; cDNA; 2458 BP. Human PRO polynucleotide #194. US2003077722-A1.

Query Match

BEREE

RESULT 54

24-APR-2003. (GETH) GENENTECH INC.

Best Local Similarity RESULT 55

Query Match

Score 2458; Pred. No. 0;

USzucz-01-MAY-2003. (GETH) GENENTECH INC. 100.0%;

Ouery Match Best Local Similarity

Best Loc RESULT 56

US-COLORDO DO COLORDO DE COLORDO

RESULT 57

ADB13259 standard; cDNA; 2458 BP. Human PRO polynucleotide #194. US2003082710-A1.

OG-MAR-2003. 06-MAR-2003. (GETH) GENENTECH INC. (GETH) GENENTECH INC. 100.0%; Score 2458;

Score 2458; Pred. No. 0;

13-MAK-2003.
1 (GETH) GENENTECH INC.
100.0%;
Query Match cimilarity 100.0%;

20-MAR-2003. (GETH) GENENTECH INC.

Best Local Similarity
RESULT 59
ID ADA12258 standard; cl
DE Human CDNA encoding FN US2003055216-A1.
PD 20-MAR-2003.
PA (GETH) GENENTECH INC

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100.0%; Score 2458; DB 9; Length 2458; 100.0%; Pred. No. 0;
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                                                                   100.0%; Score 2458; DB 9; Length 2458; 100.0%; Pred. No. 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ALMA84759 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
22003082718-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel human secreted and transmembrane protein PRO363 cDNA US2003082695-A1.
Human cDNA encoding secreted/transmembrane protein PRO363 US2003059832-A1.
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DB 9; Length 2458;

Length 2458;

DB 9;

Length 2458;

DB 9;

RESULT 76
ID ABT444
DE Human
PN US200
PD 06-FE
PA (GETF

Best Loc RESULT 75

DB 9; Length 2458;

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RESULT 69
ID ADA75
DE Human
PN US200
PD 01-MA
PA (GETH

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ACD29070 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane polypeptide cDNA #16.
US2003049633-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADB21765 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003082765-A1.
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Human secreted/transmembrane polypeptide PRO 363 cDNA.
US2003044934-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                PD 01-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 0;
                                                                                                                                                        PA (GETH) GENENTECH INC.

Query Match
100.0%; Score 2458;
Best Local Similarity 100.0%; Pred. No. 0;
RESULT 80
                                                100.0%; Score 2458; 100.0%; Pred. No. 0;
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100.0%; Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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CDNA encoding human PRO polypeptide #194
US2003082760-A1.
                                                                                  Human PRO polynucleotide #194.
US2003082702-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADA95971 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
01-MAY-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                       ADA96523 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003082690-A1.
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(GETH ) GENENTECH INC.
                13-MAR-2003.
(GETH ) GENENTECH INC.
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                                              Query Match
Best Local Similarity
RESULT 79
                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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Best Local Similarity
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US2003049817-A1.
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                                                                                                  DB 9; Length 2458;
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                                                                                                                        Human cDNA encoding secreted/transmembrane protein PRO363. US2003059780-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADA93064 standard; CDNA; 2458 BP.
Whuman CDNA encoding secreted/transmembrane protein PRO363
US200306407-A1.
27-MAR-2003.
                                                                                                                                                                                                                           DB 9;
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(GETH ) GENENTECH INC.
100.0%; Score 2458;
STY Match 100.0%; Pred. No. 0;
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24-APR-2003.
(GETH) GENENTECH INC.
ery Match 100.0%; Score 2458;
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(GETH ) GENENTECH INC.
:ry Match 100.0%; Score 2458;
:ry Match 100.0%; Pred. No. 0;
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Pred. No. 0;
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(GETH ) GENENTECH INC.
170 Match
110 Local Similarity 100.0%; Pred. No. 0;
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Pred. No. 0;
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CDNA encoding human PRO polypeptide #194.
US2003096386-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADB26832 standard; cDNA; 2458 BP.
cDNA encoding human PRO polypeptide #194.
US2003092147-A1.
                                                                                                                                                                                                                                                                                                                                                                                                               ADB25306 standard; cDNA; 2458 BP.
Human PRO polynucleotide SEQ ID NO 387.
US2003077715-A1.
24-APR-2003.
                                                                                                                                                                                                                                                                        ADA47010 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003073210-A1.
              ADA75785 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003082703-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADA93482 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003077721-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABT44573 standard; cDNA; 2458 BP. Human PR0363 cDNA. US20030027988-A1. 06-FBB-2003. (GETH ) GENENTECH INC.
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                                            US2003uor.
01-MAY-2003.
(GETH ) GENENTECH INC.
""" 100.0%;
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(GETH ) GENENTECH INC.
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RESULT 71
ID ADA47010 standard; cl
DE Human PRO polynucleof
PN US200373210-A1.
PD 17-APR-2003.
PA (GETH ) GENENTECH INC
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Best Local Similarity
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Best Local Similarity
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03-APR-2003.
(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 99
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RESULT 103
ID ABT43946 standard;
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Best Local Similarity
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Pred. No. 0;
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Human cDNA encoding secreted/transmembrane polypeptide PRO363
US2003040473-A1.
27-FBB-2003.
                  DB 9; Length 2458;
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Novel human secreted and transmembrane protein PRO363 cDNA.
US2003054516-Al.
                                                                                                                                                                  Novel human secreted and transmembrane protein PRO363 cDNA US2003082709-A1.
                                                                                                                                                                                                                                                                                                                      ADA88070 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003082700-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACH65627 standard; cDNA; 2458 BP.
Human cDNA encoding secreted/transmembrane protein PRO363
US2003044806-A1.
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24-APR-2003.
(GETH) GENENTECH INC.
(GETH) GENENTECH 100.0%; Score 2458;
ery Match 100.0%; Pred. No. 0;
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01-MAY-2003.
(GETH ) GENENTECH INC.
ery Match 100.0%; Score 2458;
               100.0%; Score 2458; 100.0%; Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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cDNA encoding human PRO polypeptide #194.
US2003082706-A1.
                                                             ADB18284 standard; cDNA; 2458 BP.
cDNA encoding human PRO polypeptide #194.
US2003077710-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADB28488 standard; cDNA; 2458 BP.
cDNA encoding human PRO polypeptide #194.
US2003082699-A1.
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31-MAX-2003.
(GETH.) GENENTECH INC.
100.0%;
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01-MAY-2003.
(GETH ) GENENTECH INC.
100.0%;
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 94
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Best Local Similarity
RESULT 95
               Query Match
Best Local Similarity
RESULT 88
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Best Local Similarity
RESULT 96
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Best Local Si
RESULT 91
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Best Le
RESULT 90
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01-MAY-2003.
(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
ery Match 100.0%; Score 2458; DB 9; Length 2458;
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                                                                                                                                                     DB 9; Length 2458;
ID ADA88622 standard; CDNA; 2458 BP.

BE NOVED human secreted and transmembrane protein PRO363 CDNA.

PN US2003073213-A1.

PD 17-APR-2003.

PA (GETH ) GENENTECH INC.

QUETY MATCh 100.0%; Score 2458; DB 9; Length 24:

Best Local Similarity 100.0%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADB22117 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003087344-A1.
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Human cDNA encoding secreted/transmembrane protein PRO363.
.US2003059782-A1.
27-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADA06791 standard; cDNA; 2458 BP.
Human secreted/transmembrane PRO polypeptide cDNA #140
US2003049638-A1.
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USDNA encoding human PRO polypeptide #194.
USD03022239-Al.
30-JAN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACD39617 standard, cDNA, 2458 BP.
3 Human cDNA encoding PR05723.
4 US2003017982-Al.
5 23-JAN-2003.
6 Query Match
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Human PRO polynucleotide #194.
US2003068793-A1.
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Human PRO polynucleotide #194.
US2003077711-A1.
                                                                                                                                                                                                                       ADA97627 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003082686-A1.
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ADB38127 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003087347-Al.
08-MAY-2003.
(GETH) GENENTECH INC.
100.0%; Score 2458; DB 9; Length 24.
ist Local Similarity 100.0%; Pred. No. 0;
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Local Similarity 100.0%;
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Best Local Similarity 100.0%;
RESULT 125
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Best Local Similarity 100.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
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   DB 9; Length 2458;
                                                                                                                                   Length 2458;
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(GETH) GENENTECH INC.
ery Match 100.0%; Score 2458; DB 9; Length 2458;
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                                                                                                                                                                  AUA92364 standard; CDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 CDNA.
US2003082712-Al.
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Novel human secreted and transmembrane protein PRO363 CDNA.
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Novel human secreted and transmembrane protein PRO363 cDNA
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Novel human secreted and transmembrane protein PRO363 CDNA.
US2003082766-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003092889-A1.
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Novel human secreted and transmembrane protein PRO363
US2003088068-A1.
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(GETH ) GENENTECH INC.
ry Match 100.0%; Score 2458;
ry Match 100.0%; Pred. No. 0;
100.0%; Score 2458; 100.0%; Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
                                              ADB23642 standard; cDNA; 2458 BP.
Human PRO polynuclectide SEQ ID NO 387.
US2003077712-A1.
24-APR-2003.
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Human PRO polynucleotide #194.
US2003087352-A1.
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22-MAY-2003.
(GETH ) GENENTECH INC.
Warch 100.0%; S
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Human PRO polymucleotide #140.
US20030554403-A1.
20-MAR-2003.
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(GETH) GENENTECH INC.
MALCh 100.0%;
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08-MAY-2003.
(GETH ) GENENTECH INC.
100.0%;
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DE Novel human secreted at
PD US2003082712-A1.
PD 01-MAY-2003.
PA (GETH ) GENEWTECH INC.
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(GETH ) GENENTECH INC.
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          Best Local Similarity
RESULT 107
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RESULT 111
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Best Local Similarity
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Length 2458;
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Pred. No. 0;
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Length 2458;
                                                ADB66599 standard, cDNA, 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003082764-Al.
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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                                                                                                                                                                                        ADB84996 standard; cDNA; 2458 BP.
Human PRO polynucleotide #65.
US2003073817-A1.
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Human PRO polynucleotide #194.
US2003082698-A1.
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Human PRO polynucleotide #194.
US2003082762-A1.
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Human PRO polynucleotide SEQ ID NO 387
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        Human PRO polynucleotide #65.

US2003092890-A1.

US-2003092890-A1.

(GETH ) GENENTECH INC.

ETY MATCH

SCORE 2458; DB 10; Length 2458; Bt Local Similarity 100.0%; Pred. No. 0;
                                                                                                                                                                                                                                                    ADB83865 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003069397-A1.
US2003069397-A1.
(GPAR-2003.
(GPAH ) GENENTECH INC.
100.0%; Score 2458; DB 10; Length 2458;
st Local Similarity 100.0%; Pred. No. 0;
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(GETH) GENENTECH INC.
ery Match 100.0%; Score 2458; DB 10; Length 2458;
ery match 100.0%; Pred. No. 0;
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003082696-Al.
                                                                                                                        ADB47135 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003082687-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003092887-Al.
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24-APR-2003.
(GETH) 100.0%; SCORE 2458;
ATV MATCh 100.0%; Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Human PRO polynucleotide SEQ ID NO 387.
US2003077717-A1.
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Human PRO polynucleotide sequence #16.
US2003083248-A1.
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US2003082697-A1.
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ADB84750 standard; cDNA; 2458 BP.
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01-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%;
RESULT 134
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Best Local Similarity
RESULT 126
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Best Local Similarity
RESULT 127
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RESULT 128
ID ADB86742 standard;
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Best Local Similarity
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Query Match 100.0%; Score 2458; DB 10; Length 2458; Best Local Similarity 100.0%; Pred. No. 0; RESULT 136
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ADC43706 standard; cDNA; 2458 BP.
Human cDNA encoding secreted/transmembrane protein, PRO363.
US2003054986-A1.
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Human cDNA encoding secreted/transmembrane protein, PRO363
US2003054405-Al.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003082692-Al.
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Human cDNA encoding secreted/transmembrane protein, PRO363
US2003049684-A1.
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Human cDNA encoding secreted/transmembrane protein PRO363
US2003049681-A1.
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(GETH) GENENTECH INC.
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(ery Match 100.0%; Score 2458; DB 10;
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20-MAR-2003.
(GETH ) GENENTECH INC.
(BLY MATCh 100.0%; Score 2458;
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Pred. No. 0;
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Human PRO polynucleotide #140.
US2003027754-A1.
06-FEB-2003.
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(CFTH) GENENTECH INC.
100-17 Match
Best Local Similarity 100.0%;
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(GETH ) GENENTECH
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ADC36858 standard; cDNA; 2458 BP.
Human PRO polynucleotide #65.
US2003088065-A1.
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Human cDNA encoding secreted/transmembrane protein, PRO363
US2003064407-Al.
                                                          PRO363
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Human cDNA encoding secreted/transmembrane protein PRO363.
102003069403-A1.
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Human cDNA encoding secreted/transmembrane protein,
122003073131-Al.
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Human cDNA encoding secreted/transmembrane protein,
US2003069178-A1.
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Human cDNA encoding secreted/transmembrane protein,
US2003072745-A1.
                                                         Human cDNA encoding secreted/transmembrane protein, US2003060406-A1.
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Human cDNA encoding secreted/transmembrane protein,
US20031068648-A1.
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No. 0;
 100.0%; Score 2458; 100.0%; Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred.
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Human PRO polynucleotide #140.
US2003064375-A1.
                                            CDNA; 2458 BP
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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          Best Local Similarity RESULT 144
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                                            ADC66530 standard;
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Novel human secreted and transmembrane protein PRO363 cDNA.
US2003092106-Al.
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ADC49879 standard, cDNA, 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003092107-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003092105-A1.
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Novel human secreted and transmembrane protein PRO363
                                                                                            ADC41723 standard; cDNA; 2458 BP.
Human cDNA encoding secreted/transmembrane protein,
US2003104998-Al.
                                               DB 10;
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Pred. No. 0;
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08-MAY-2003.
(GETH ) GENENTECH INC.
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                                              Query Match
Best Local Similarity
RESULT 154
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RESULT 156
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(GETH ) GENENTECH
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Length 2458;

Length 2458;

DB 10;

Length 2458

DB 10;

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ADC48512 standard; cDNA; 2458 BP.

Human PRO polynucleotide #194.

1 US2003194773-A1.

1 GETH ) GENENTECH INC.

Query Match

A GOOT AD A CONTROL OF CONTRO
                                                                                                                         AUCSB538 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein cDNA Seq ID387.
US2003087346-Al.
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Novel human secreted and transmembrane protein PRO363 CDNA.
US2003092104-Al.
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Novel human secreted and transmembrane protein PRO363 cDNA.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003082546-A1.
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Novel human secreted and transmembrane protein PRO363
US2003105288-A1.
(GETH ) GENENTECH INC.
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Novel human secreted and transmembrane protein PRO363
US2003087348-A1.
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Best Local Similarity 100.0%; Pred. No. 0
RESULT 179
ID ADC69623 standard; CDNA; 2458 BP.
DE CDNA encoding human PRO polypeptide #194.
PN US2003194770-A1.
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Human PRO polynucleotide #140.
US2003083461-A1.
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(GETH ) GENENTECH INC.

Query Match 100.0%;

Best Local Similarity 100.0%;
                     08-MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                                                               Best Local Similarity
                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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Best Local Similarity
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Best Local Similarity
RESULT 178
US2003087360-A1.
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Novel human secreted and transmembrane protein cDNA Seg ID387.
US200308736-A1.
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Novel human secreted and transmembrane protein cDNA Seg ID387.
US2003087364-A1.
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Novel human secreted and transmembrane protein cDNA Seg ID387.
US2003087363-A1.
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Novel human secreted and transmembrane protein cDNA Seq ID387
US2003087395-A1.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                 Novel human secreted and transmembrane protein cDNA Seq ID387
US2003087365-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA, US2003087367-Al.
08-MAY-2003.
(GETH ) GENENTECH INC.
100.0%; Score 2458; DB 10; Length; St Local Similarity 100.0%; Pred. No. 0;
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Novel human secreted and transmembrane protein PRO363 cDNA
082003087361-A1.
08-MAY-2003.
(GETH ) GENENTECH INC.
100.0%; Score 2458; DB 10; Length
st Local Similarity 100.0%; Pred. No. 0;
Novel human secreted and transmembrane protein PRO363 cDNA
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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                                                                                                                                                                            CDNA; 2458 BP
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(GETH ) GENENTECH INC.

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08-MAY-2003.
(GETH ) GENENTECH INC.
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08-MAY-2003.
(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 169
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Best Local Similarity
RESULT 171
                                                                                                                 Best Local Similarity RESULT 163
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Best Local Similarity
RESULT 165
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(GETH ) GEN
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Length 2458;

Length 2458

CDNA

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ADC47960 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003194771-A1.
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100.0%;
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                                                                                               DB 10; Length 2458;
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Novel human secreted and transmembrane protein PRO363 CDNA.
                                                                                                                                     ADD07801 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2002193299-A1.
                                                                                                                                                                                                                                          ADC78076 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003096972-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003087354-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003092103-A1.
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Human secreted/transmembrane PRO polypeptide cDNA
US2003105011-A1.
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                       ADD10041 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003194776-A1.
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05-JUN-2003.
(GETH ) GENENTECH INC.
100.0%; S/
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08-MAY-2003.
(GETH ) GENENTECH INC.
100.0%;
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12-MAY-2003.
(GETT.) GENENTECH INC.
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17-APR-2003.
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16-OCT-2003.
(GETH ) GENENTECH INC. 100.0%;
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US2003194774-A1.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 184
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 Best Local Similarity
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RESULT 183
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RESULT 190
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Novel human secreted and transmembrane protein PRO363 CDNA.
US2003105291-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADD41202 standard; cDNA; 2458 BP. Novel human secreted and transmembrane protein PRO363 cDNA US2003203438-A1.
                                                                                                                                                                          ADC77830 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003088066-A1.
                                                                                                                                                                                                                                                                                                                    ADC80020 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA US2003087358-A1.
                                                     ADD08872 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADD07121 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein, PRO363 cDNA
US2002193300-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human secreted/transmembrane PRO polypeptide cDNA #32
US2003105013-A1.
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Pred. No. 0;
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Pred. No. 0;
   DB 10;
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Score 2458;
Pred. No. 0;
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Pred. No. 0;
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                                                                                                                         Score 2458;
Pred. No. 0;
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Pred. No. 0;
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US2003059783-A1.
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Human PRO polynucleotide #194.
US2003194775-A1.
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                                                                                                                         100.08;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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ID ADC83368 standard;
                                                   ADD08872 standard;
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Best Local Similarity 100.0%; Pred. No. 0; RESULT 209
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Human PRO polynucleotide #194.
US2003203431-A1.
30-CCT-2003.
(GETH ) GENENTECH INC.
ery Match
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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 2458; DB 10; Length 2458;
ery Match 100.0%; Pred. No. 0;
                                          DB 10; Length 2458;
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Novel human secreted and transmembrane protein PRO363 CDNA.
                                                                                                                                                                                                                                                 ADD51039 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003105290-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADD37105 standard; cDNA; 2458 BP.
Human secreted/transmembrane PRO polypeptide cDNA #32.
US2003105012-A1.
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(GETH ) GENENTECH INC.

(GETH ) GENENTECH INC.

100.0%; Score 2458;

Lery Match 100.0%; Pred. No. 0;
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16-OCT-2003.
(GETH ) GENENTECH INC.
100.0%; Score 2458;
iery Match 100.0%; Pred. No. 0;
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(GETH) GENENTECH INC.
(GETY Match 100.0%; Score 2458;
                 PA (GETH ) GENENTECH INC.

Query Match
Best Local Similarity 100.0%; Pred. No. 0;
RESULT 200
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                                                                                                                                                                                             Score 2458;
Pred. No. 0;
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Pred. No. 0;
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cDNA encoding human PRO polypeptide #194.
US2003194779-A1.
                                                                                             ADD52341 standard; cDNA; 2458 BP.

CDNA encoding human PRO polypeptide #194.
US2001194769-A1.
16-OCT-2003.
(GETH) GENENTECH INC.
EXT MATCh

SECOLO SIMILARITY 100.0%; Pred. No. 0%
                                                                                                                                                                                                                                                                                                                                                                                                                     cDNA encoding human PRO polypeptide #194.
US2003194792-A1.
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Human PRO polynucleotide #140.
US2003077594-A1.
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Human PRO polynucleotide #140.
US2003077593-A1.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 204
ID ADD55475 standard; c
DE Human PRO polynucleo
PN US2003077593-A1.
PD 24-APR-2003.
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Best Local Similarity
RESULT 202
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       30-OCT-2003
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Best Local S
RESULT 207
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100.0%; Score 2458; DB 10; Length 2458; 100.0%; Pred. No. 0;
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Novel human secreted and transmembrane protein PRO363 cDNA, US2003105289-A1.
05-JUN-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                     ADD54204 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003203432-Al.
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Human cDNA encoding secreted/transmembrane protein, PRO363
US2003096744-Al.
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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                                                                                                                        Human PRO polynucleotide #194.
US2003203430-A1.
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Human PRO polynucleotide #140.
US2002132253-A1.
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Human PRO polynucleotide #65.
US2003096970-A1.
ADD50520 standard, cDNA, 2458 BP.
Human PRO polynucleotide #65.
US2003096971-A1.
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PA (GETH) GENENTECH INC.
Query Match 100.0%;
Best Local Similarity 100.0%;
RESULT 216
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(GETH ) GENENTECH INC.
                                                 ) 22-MAY-2003.
1 (GETH ) GENENTECH INC.
Query Match 1
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
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                                                                                               Best Local Similarity RESULT 210
                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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Length 2458;

Length 2458;

Length 2458;

RESULT 218
ID ADE040
DE Human
PN US2003
PD 23-OCT
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100.0%; Score 2458; DB 10; Length 2458; 100.0%; Pred. No. 0;
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003194791-Al.
                                                                                                                                                                                                                                                                                                                                                                               ADE33432 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003194767-A1.
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Human cDNA encoding secreted/transmembrane protein, PRO363
US2003194781-A1.
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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cDNA encoding human PRO polypeptide #194.
US2003201417-Al.
                                                                                                ADE17837 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003199023-A1.
23-OCT-2003.
(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
est Local Similarity 100.0%; Pred.
                                                                                                                                                                                                                                            ADD91969 standard; cDNA; 2458 BP.
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US2003199025-A1.
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DRE18941 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003199026-Al.
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16-OCT-2003.
(GETH ) GENENTECH INC.
100.0%; S
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US2003199053-A1.
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PA (GETH ) GENENTECH INC.

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BESE LOCAL Similarity 100.0%;

RESULT 230
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
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Best Local Similarity
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US2003194772-A1.
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RESULT 234
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Novel human secreted and transmembrane protein PRO363 cDNA.
US2003087304-A1.
08-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                               ADE32328 standard; cDNA, 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
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Human cDNA encoding secreted/transmembrane protein, PRO363
082003203435-A1.
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Human cDNA encoding secreted/transmembrane protein, PRO363
US2003203414-A1.
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                                                                                                                                                 ADE31890 standard; cDNA; 2458 BP.
Human cDNA encoding secreted/transmembrane protein PRO363.
US200306847-A1.
10-APR-2003.
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(GETH ) GENENTECH INC.
:ry Match 100.0%; Score 2458;
rry Match 100.0%; Pred. No. 0;
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(GETH ) GENENTECH INC.
:ry Match 100.0%; Score 2458;
rry Match 100.0%; Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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cDNA encoding human PRO polypeptide #194.
US2003203428-A1.
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US2003199056-A1.
23-OCT-2003.
          ADE04031 standard; cDNA; 2458 BP.
Human PRO Polynucleotide #194.
23-0CT-2003.
GETH J GENENTECH INC.
(GETH J GENENTECH INC.
LUCY MALCH 100.0%; SCORE
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Human PRO polynucleotide #194.
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30-OCT-2003.
(GETH ) GENENTECH INC.
100.0%;
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(GETH ) GENENTECH INC.
100.0%;
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 222
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Best Local Similarity
RESULT 226
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Best Local Similarity
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Best Local Si
RESULT 224
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RESULT 223

RESULT 225

Length 2458;

Length 2458;

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Query Match

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GENENTECH INC.
h 100.0%; Score 2458; DB 10; Length 2458;
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100.0%; Score 2458; DB 10; Length 2458; 100.0%; Pred. No. 0;
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Human cDNA encoding secreted/transmembrane protein, PRO363
125003195333-A1.
16-CGT-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADG21237 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA US2003207355-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003207384-A1.
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Local Similarity 100.0%; Pred. No. 0;
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(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
100.0%; Score 2458;
                                                  ADD89616 standard; cDNA; 2458 BP. Human PRO polynucleotide #194. 0202003199028-Al. 23-OCT-2003. (GETH ) GENENTECH INC.
                                                                                                                                                                                                      ADE40900 standard; cDNA; 2458 BP. Human PRO polynucleotide #194. 032003199031-A1. 03-0CT-2003. (GETH ) GENENTECH INC.
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Human PRO polynucleotide #194.
US2003199034-A1.
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Human PRO polynucleotide #194.
US2003207370-A1.
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Best Local Similarity 100.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 248
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                 Best Local Similarity RESULT 246
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(GETH ) GEN
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(GETH ) GEN
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                                        DB 10; Length 2458;
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Human cDNA encoding secreted/transmembrane protein, PRO363.
US2003203433-Al.
30-OCT-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADE26492 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003087305-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003194766-A1.
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16-OCT-2003.
(GETH ) GENENTECH INC.
(GETH) GENENTECH INC. 0%; Score 2458;
ery Match 100.0%; Pred. No. 0;
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(GETH ) GENENTECH INC.

(GETH ) ABLCh

100.0%; Score 2458;
         PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Fred. No. 0;
RESULT 237
                                                                                                                                                                                                                                             ADD95926 standard; CDNA; 2458 BP.
Human PRO polymucleotide #194.
23-OCT-2003.
(GETH ) GENENTECH INC.
100.04; Score 2458;
EL Local Similarity 100.04; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                    DE CDNA encoding human PRO polypeptide #194.
PN US2003199064-Al.
PD 23-OCT-2003.
PA (GETH) GENENTECH INC.
Query Match
100.0%; Score 2458;
Best Local Similarity 100.0%; Fred. No. 0,
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Query Match
100.0%; Score 2458;
Best Local Similarity 100.0%; Pred. No. 0;
RESULT 245
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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cDNA encoding human PRO polypeptide #194.
US2003203429-A1.
30-OCT-2003.
(GETH ) GENENTECH INC.
PLY MAtch
LLOCAL Similarity 100.0%; Score 2458;
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cDNA encoding human PRO polypeptide #194.
US2003207418-Al.
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Human PRO polynucleotide #194.
US2003199032-A1.
                                                                                           ADE43137 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003199033-A1.
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RESULT 239
ID ADE22812 standard; CDNA; 2458 BP.
                                                                                                                            100.0%;
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(GETH ) GENENTECH INC.
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RESULT 242
ID ADE32880 standard; cl
DE Novel human secreted
PD US2003194766-A1.
PD 16-OCT-2003.
PA (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
RESULT 241
ID ADE26492 standard; cl
DE Novel human secreted
PN US2003087305-A1.
PD 08-MAY-2003.
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RESULT 243
                                                                                                                                                                                         Query Match
Best Local Similarity
RESULT 238
   23-OCT-2003
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Query Match

Page 15

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ADI63536 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003207387-A1.
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Best Local Similarity
RESULT 269
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Best Local Similarity
RESULT 271
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Human cDNA encoding secreted/transmembrane protein, PRO363.
US2003206915-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA.
US2003207379-A1.
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Novel human secreted and transmembrane protein PRO363 CDNA, 22003207381-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003207385-A1.
                                                                                                                                                        ADG52655 standard; cDNA; 2458 BP.
Human cDNA encoding secreted/transmembrane protein,
US2003216561-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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US2003050457-A1.
                              ADG80577 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003207373-A1.
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Human PRO polynucleotide #194.
US2003207372-A1.
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Novel human secreted and transmembrane protein PRO363 CDNA.
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Novel human secreted and transmembrane protein PRO363 cDNA.
US2003207377-A1.
                                                                                                              ADH81950 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003207388-Al.
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Novel human secreted and transmembrane protein PRO363 cDNA
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Human cDNA encoding secreted/transmembrane protein PRO363.
US2003032023-A1.
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Novel human secreted or transmembrane protein PRO5723 DNA
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US2002160384-A1.
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CDNA encoding human PRO polypeptide #65.
20-FEB-2003.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003100722-A1.
                                      Best Local Similarity 100.0%; Pred. No. 0; RESULT 283

ID AD165037 standard; cDNA; 2458 BP.

DE Novel human secreted and transmembrane protein PRO363 cDNA PD 06-NOV-2003.
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Novel human secreted and transmembrane protein PRO363 cDNA
15.009288-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA US2003100735-Al.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003100717-A1.
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US2003100087-A1.
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29-Mav.nn.
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Novel human secreted and transmembrane protein PRO363 cDNA.
US2003087357-Al.
                                                                                                     ACD42474 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003050239-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003087353-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA.
US2003087385-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003045687-A1.
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Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cDNA encoding human PRO polypeptide #194.
US2003004311-A1.
                                                                                                                                                                                                                                                                                                                                                                           ABX64291 standard; cDNA, 2458 BP.
cDNA encoding human PRO363 polypeptide.
025002103125-A1.
01-AUG-2002.
GETH ) GENENTECH LTD.
CRY MATCH
TY LOOAL Similarity 100.0%; Score 245
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACA67187 standard; cDNA; 2458 BP
(GETH ) GENENTECH INC.
100.0%; S.
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02-JAN-2003.
(GETH ) GENENTECH INC.
100.0%; SC
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08-MAY-2003.
(GETH ) GENENTECH INC.
100.0%;
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06-MAR-2003.
(GETH ) GENENTECH INC.
Match '...tev 100.0%;
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(GETH ) GENENTECH INC.
MATCh '1-rity 100.0%;
                                                                                                                                                 PD 13-MAR-2003.
PA (GETH ) GENENTECH INC.
QUERY MATCh 100.0%;
Best Local Similarity 100.0%;
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08-MAY-2003.
(GETH ) GENENTECH INC.
MACCh ''THICK 100.0%;
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(GETH ) GENENTECH INC.
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RESULT 278
ID ADM82567 standard, cI
DE Novel human secreted
PN US2003087355-A1.
PD 08-MXY-2003.
PA (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity RESULT 277
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Best Local Similarity
RESULT 279
                                                               Best Local Similarity RESULT 274
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Best Local Similarity
RESULT 276
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08-MAY-2003.
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Query Match

Query Match

Query Match

Query Match

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AUD78426 standard; CDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 CDNA.
US2003100737-A1.
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Best Local Similarity 100.0%;
RESULT 310
                                                                                                                Best_Local Similarity RESULT 302
                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
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                                                                                                         Query Match
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                                                                                                                                                                                                                                                                                                        ADE39054 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003096332-A1.
22-MAY-2003.
(GETH) GENENTECH INC.
PRY MATCH
OCAL Similarity 100.0%; Score 2458; DB 12; Length 2458; Local Similarity 100.0%; Pred. No. 0;
                                                                                                                       Length 2458;
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                                   ADD86880 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003100738-A1.
                                                                                                                                                                    ADE20757 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003100734-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADE48392 standard; cDNA; 2458 BP.
Human cDNA encoding secreted/transmembrane protein, PRO363.
US2003104536-A1.
05-JUN-2003.
(GETH ) GENENTECH INC.
                                                                                                                       DB 12;
                                                                                29-MAY-2003.
(GETH ) GENENTECH INC.
ery Match 100.0%; Score 2458;
et Local Similarity 100.0%; Pred. No. 0;
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29-MAY-2003.
(GETH ) GENENTECH INC.
ery Match 100.0%; Score 2458;
c'milarity 100.0%; Pred. No. 0;
Best Local Similarity 100.0%; Fred. No. 0; RESULT 292
                                                                                                                                                                                                                                                          Score 2458;
Pred. No. 0;
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(GETH ) GENENTECH INC.

(ery Match 100.0%; Fred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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US2003100711-A1.
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Human PRO polynucleotide #65.
US2003100727-A1.
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Human PRO polynucleotide #194.
US2003211571-A1.
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29-MAY-2003.
(GETH ) GENENTECH INC.
100.0%; Sr
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PA (GETH) GENENTECH INC.
Query Match 100.0%;
Best Local Similarity 100.0%;
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29-MAY-2003.
(GETH ) GENENTECH INC.
MATCh 100.0%;
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(GETH ) GENENTECH INC.
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RESULT 294
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RESULT 296
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Best Local Similarity
RESULT 298
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RESULT 299
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DB 12; Length 2458;
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                                                                                                                                                                                                                                                                                                                                                                                      ADE21249 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003100736-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel human secreted and transmembrane protein PRO363 cDNA US2003100733-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADD77364 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003100732-A1.
                                                                                            ADE41313 standard; cDNA; 2458 BP.
Human secreted/transmembrane PRO polypeptide cDNA #32.
25203100497-A1.
29-MAY-2003.
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Best Local Similarity 100.0%; Pred. No. 0;
RESULT 307
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Pred. No. 0;
                                           DB 12;
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GBTH) GENENTECH INC.
(GETH) GENENTECH INC.

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100.0%; Score 2458;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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cDNA encoding human PRO polypeptide #194.
US2003092108-A1.
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US2003100708-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADD75576 standard; cDNA; 2458 BP.
Human PRO polynucleotide #65.
US2003100064-A1.
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Best Local Similarity 100.0%;
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(GETH ) GENENTECH INC.
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Query Match
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Human PRO polynucleotide #194.
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PA (GETH ) GENENTECH INC.

Query Match

100.0%; Score 2458; DB 12; Length 2458;
Best Local Similarity 100.0%; Pred. No. 0;

RESULT 313

ID ADE24559 standard; CDNA; 2458 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 12; Length 2458;
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           Novel human secreted and transmembrane protein PRO363 cDNA. US2003100718-A1.
29-MAY-2003.
(GETH ) GENENTECH INC.
100.0%; Score 2458; DB 12; Length 2 t Local Similarity 100.0%; Pred. No. 0;
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Novel human secreted and transmembrane protein PRO363 cDNA.
US2003100719-Al.
                                                                                                           ADD85560 standard; CDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 CDNA
US2003100721-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADD76866 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003100715-A1.
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PA (GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 0;
RESULT 314
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PA (GETH ) GENENTECH INC.

Query March 100.0%; Score 2458;

Best Local Similarity 100.0%; Pred. No. 0;

RESULT 315
                                                                                                                                                PD 29-MAY-2003.
PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%; Pred. No. 0;
RESULT 312
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Pred. No. 0;
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Pred. No. 0;
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Local Similarity 100.0%; Pred. No. 0;
                                                                                                                                                                                                                            ADE23916 standard; cDNA; 2458 BP.
cDNA encoding human PRO polypeptide #194.
US2003092110-A1.
                                                                                                                                                                                                                                                                                                                                                           cDNA encoding human PRO polypeptide #194.
US2003092111-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADE05109 standard; cDNA; 2458 BP.
Human PRO polynucleotide #65.
US2003100726-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADD87384 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003203439-A1.
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 ADD76068 standard; cDNA; 2458 BP
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29-MAY-2003.
(GETH ) GENENTECH INC,
100.0%;
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PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 100.0%;
RESULT 319
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Best Local Similarity
RESULT 316
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Best Local Si
RESULT 318
                                                                       Query Match
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Score 2458; DB 12; Length 2458; Pred. No. 0;
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                                Length 2458;
                                                                                                                                                        'Match 100.0%; Score 2458; DB 12; Length 2458;
Local Similarity 100.0%; Pred. No. 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human cDNA encoding secreted/transmembrane protein, PRO363.
US2003130181-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AUD77610 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
                                                                           ADD78102 standard, cDNA, 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003100731-Al.
                   Luciy Match Best Local Similarity 100.0%; Score 2458; DB 12; Best Local Similarity 100.0%; Pred. No. 0; RESULT 320
                                                                                                                                                                                                                                                                                     DB 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 12;
                                                                                                                                                                                                                                                     16-OCT-2003.
A (GETH) GENENTECH INC.
100.0%; Score 2458;
Best Local Similarity 100.0%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 2458;
Pred. No. 0;
                                                                                                                                                                                                    ADE18389 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003194794-A1.
                                                                                                                                                                                                                                                                                                                               ADE88698 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003199054-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE89493 standard; cDNA; 2458 BP.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
(GETH ) GENENTECH INC.
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GERRITSEN M E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (TUMA/) TUMAS D.
(WILL/) WILLIAMS P M.
(WOOD/) WOOD W I.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EATON D L.
FERRARA N.
FILVAROFF E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GODOWSKI P J. GIRMALDI J C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             KLJAVIN I J.
KUO S S.
NAPIER M A.
PAN J.
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ROY M A.
SHELTON D L.
                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best_Local Similarity RESULT 325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BOTSTEIN D.
DESNOYERS L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GURNEY A L. HILLAN K J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GODDARD A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BAKER K P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FONG S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GAO W.
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EATO/)
FERR/)
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GERB/)
GERR/)
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Human cDNA encoding secreted/transmembrane protein, PRO363
                   .S2003150...
23-0CT-2003.
(GETH ) GENENTECH INC.
100.0%;
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                                                                        Query Match
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                                                ) 29-MAY-2003.

A (GETH) GENENTECH INC.

Query Match 100.0%; Score 2458; DB 12; Length 2458;

Best Local Similarity 100.0%; Pred. No. 0;
                                                                                                                                                                                                                       Score 2458; DB 12; Length 2458;
Pred. No. 0;
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ADD77856 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003100730-A1.
                                                                                                                                 ADD85314 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADD77112 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003100716-A1.
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Human cDNA encoding secreted/transmembrane protein, PRO363
22003195345-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003100720-A1.
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16-COT-2003.
(GETH ) GENENTECH INC.
ELY MATCH 100.0%; SCOTE 2458;
Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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US2003100710-A1.
29-MAY-2003.
                                                                                                                                                                                                                                                                                                                                                                                                              ADD74584 standard; cDNA; 2458 BP.
Human PRO polynucleotide #65.
US2003100713-A1.
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25-MAV-----
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Human PRO polynucleotide #65.
US2003100724-A1.
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29-MAX-2003.
(GETH ) GENENTECH INC.
100.0%; Sr
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) GENENTECH INC.
100.0%; S
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29-MAY-2003.
(GETH ) GENENTECH INC.
100.0%;
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29-MAY-2003.
(GETH) GENENTECH INC.
Match '...tt 100.0%;
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(GETH ) GENENTECH INC.
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Best Local Similarity
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RESULT 3
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Length 2458;
   Length 2458;
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                                                                 Human cDNA encoding secreted/transmembrane protein, PRO363 US2003195148-A1.
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     12;
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ry Match 100.0%; Score 2458;

r Accal Similarity 100.0%; Pred. No. 0;
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(GETH ) GENENTECH INC.
ery Match 100.0%; Score 2458;
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Score 2458;
Pred. No. 0;
                                                                                                                                       Score 2458;
Pred. No. 0;
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Pred. No. 0;
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                                                                                                                                                                                     ADE94718 standard; cDNA; 2458 BP.
cDNA encoding human PRO polypeptide #194.
US2003199027-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cDNA encoding human PRO polypeptide #194
                                                                                                                                                                                                                                                                                                                                                                                                                                              cDNA; 2458 BP. cDNA; 2458 BP. cDNA encoding human PRO363 polypeptide. US2003194760-A1.
                                                                                                                                                                                                                                                                                                     RESULT 337

ID ADE91129 standard; cDNA; 2458 BP.
DE Human PRO polynucleotide #194.
PN US2003199061-Al.
                                                    ADF45621 standard; cDNA; 2458 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADE93380 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003199060-A1.
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                                                                                                  PD 16-OCT-2003.

PA (GETH) GENENTECH INC.

QUETY MATCh

BEST LOCAL SIMILARITY 100.0%;

RESULT 336
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                     Best Local Similarity
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              Best Local Similarity
RESULT 335
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RESULT 338
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADE95270 standard;
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(GETH ) GENENTECH INC.
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                                        Query Match
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RESULT 354
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RESULT 361
                                                                                                                                                                                                                                                                ADF26843 standard; 100.0%; Score 2458; DB 12; Length 24:

ADF26843 standard; CDNA; 2458 BP.

BB Human CDNA encoding secreted/transmembrane protein, PRO363.

PN 02200319436-A1.

PD 23-CCT-2003.

PA (GETH ) GENENTECH INC.

Query Match

Best Local Similarity 100.0%; Score 2458; DB 12; Length 7.7

ID ADF27479 standard; CDNA; 2458 BP

PR UN22003199437-A1.

PD 23-CCT-20^7

PA (CT-20^7)
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PA (GETH ) GENENTECH INC.

Query Match 100.0%; Score 2458; DB 12; Length 2458;

Best Local Similarity 100.0%; Pred. No. 0;

RESULT 349

ID ADE90577 standard; CDNA; 2458 BP.
                                                          Length 2458;
                                                                                                                                                                                                                                                                                                                                             Length 2458;
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Whuman CDNA encoding secreted/transmembrane protein, PRO363.
US2003211092-A1.
13-NOV-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADF41073 standard; cDNA; 2458 BP.
Muman cDNA encoding secreted/transmembrane protein, PRO363.
US2003199435-Al.
23-OCT-2003.
(GETH) GENENTECH INC.
100.0%; Score 2458; DB 12; Length 2
Et Docal Similarity 100.0%; Pred. No. 0;
                                                                                                         ADF33376 standard; cDNA; 2458 BP.

Human cDNA encoding secreted/transmembrane protein, PRO363.
US2003194780-Al.
16-OCT-2003.
(GETH ) GENEWTECH INC.
100.0%; Score 2458; DB 12; Length ist Local Similarity 100.0%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADE92276 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003199051-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADF32752 standard; cDNA; 2458 BP.
Human cDNA encoding secreted/transmembrane protein, PRO363
US2003211091-A1.
                                                         DB 12;
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                                                                                                                                                                                                                                                   CDNA encoding human PRO polypeptide #194. US2003199029-A1. 23-OCT-2003. (GETH) GENENTECH INC. 100.0%; Score 2458; Bt Local Similarity 100.0%; Pred. No. 0;
                                                         Score 2458;
Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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US2003199063-A1.
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13-NOV-2003.
(GETH ) GENENTECH INC.
100.0%;
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(GETH ) GENENTECH INC.
(ALCh ) GENENTECH INC.
                       PA (GETH ) GENENTECH INC.

Query Match
Best Local Similarity 100.0%;
RESULT 344
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PA (GETH ) GENENTECH INC.

QUENY MARCh

BEST LOCAL Similarity 100.0%;

RESULT 348
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2003.
.tH ) GENENTECH .
.rY Match
Best Local Similarity
RESULT 350
ID ADF41073 stand*
DE Human CDNA
PN US20031*
PD 23-
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Best Local Similarity
RESULT 351
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Best Local Similarity
RESULT 352
                                                                                                                                                                                                   Query Match
Best Local Similarity
RESULT 345
PD 30-OCT-2003.
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100.0%; Score 2458; DB 12; Length 2458; 100.0%; Pred. No. 0;
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Novel human secreted and transmembrane protein PRO363 cDNA.
                                                                                                                                                                                                                                                                                                                                                           ADF46245 standard; cDNA; 2458 BP.

Human cDNA encoding secreted/transmembrane protein, PRO363
182003195344-A1.
16-0CT-2003.
(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADG05642 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA US2003096959-A1.
                                                      ADF26219 standard; cDNA; 2458 BP.
Human cDNA encoding secreted/transmembrane protein, PRO363
US2003199674-Al.
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Human cDNA encoding secreted/transmembrane protein, PRO363
US2003194410-A1.
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100.0%; Score 2458; DB 12; 100.0%; Pred. No. 0;
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Best Local Similarity 100.0%; Pred. No. 0;
RESULT 356
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100.0%; Pred. No. 0;
                                                                                                                                                    100.0%; Score 2458; 100.0%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                         Score 2458;
Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Human PRO polynucleotide #65.
US2003096962-A1.
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Human PRO polynucleotide #194.
US2003207352-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDNA; 2458 BP.
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(GETH) GENENTECH INC.
Query Match
Best Local Similarity 100.0%;
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(GETH ) GENENTECH INC.
                                                                                                               23-OCT-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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               Best Local Similarity RESULT 353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADG11878 standard;
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Best Local Similarity
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100.0%; Score 2458; DB 12; Length 2458;
100.0%; Score 2458; DB.12; Length 2458; 100.0%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                   Length 2458;
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                                                                                                                                                                                                                                                                                                                                                     ADG24282 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003207426-A1.
                                                                                                                                                   DB 12;
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(GETH ) GENENTECH INC.
17 Match 100.0%; Score 2458;
TY Match 100.0%; Pred. No. 0;
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06-NOV-2003.
(GETH ) GENENTECH INC.
(GETH ) GENENTECH 100.0%; Score 2458;
ery Match 100.0%; Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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A (GETH ) GENENTECH INC.
100.0%; SCORE 2458;
Best Local Similarity 100.0%; Pred. No. 0;
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                                                    ADG20159 standard, cDNA, 2458 BP.
CDNA encoding human PRO polypeptide #194.
US2003207376-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADGI6773 standard; cDNA; 2458 BP.
cDNA encoding human PRO polypeptide #194.
US2003207359-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human PRO polynucleotide #194.
US2003208055-Al.
                                                                                                                                                                                              ADF98065 standard; cDNA; 2458 BP.
Human PRO polynucleotide #194.
US2003207422-A1.
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Human PRO polynucleotide #194.
US2003207351-A1.
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Human PRO polynucleotide #194.
US2003207353-A1.
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ry Match
t Local Similarity 100.0%;
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Human PRO polynucleotide #194.
US2003207375-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
               Best Local Similarity RESULT 362
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Best Local Similarity
RESULT 367
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DB 12; Length 2458;
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                                     ADG11259 standard; cDNA; 2458 BP.

Novel human secreted and transmembrane protein PRO363 cDNA.
US2003096667-A1.
22-MAY-2003.
                                                                                                                                                                                                                                                                                                                                           ADG08393 standard, cDNA, 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003207424-A1.
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Novel human secreted and transmembrane protein PRO363 CDNA.
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Novel human secreted and transmembrane protein PRO363
US2003207389-Al.
                                                                                                                                                                                                                                                                                         DB 12;
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(GETH ) GENENTECH INC.
100.0%; Score 2458;
TY Match 100.0%; Pred. No. 0;
                                                                                                       (GETH) GENENTECH INC.

ry Match 100.0%; Score 2458; t Local Similarity 100.0%; Pred. No. 0;
   100.0%; Pred. No. 0;
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100.0%; Pred. No. 0;
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Pred. No. 0;
                                                                                                                                                                                      ADG13336 standard; cDNA; 2458 BP.
cDNA encoding human PRO polypeptide #194.
US2003207357-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADG15563 standard; cDNA; 2458 BP.
cDNA encoding human PRO polypeptide #194.
US2003219885-A1.
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Human PRO polynucleotide #194.
US2003207423-A1.
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US2003207374-A1.
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PA (GETH) GENENTECH INC.
Query Match 100.0%;
Best Local Similarity 100.0%;
RESULT 378
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC
. Best Local Similarity RESULT 371
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                                                                                                                                              Best Local Similarity RESULT 372
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ADG57472 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003207362-A1.
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                                            06-NOV-2003.
(GETH ) GENENTECH INC.
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RESULT 398
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               ADG24920 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA.
US2003207437-Al.
06-NOV-2003.
(EGTH) GENENTECH INC.
1100.0%; Score 2458; DB 12; Length 2
EL Local Similarity 100.0%; Pred. No. 0;
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Novel human secreted and transmembrane protein PRO363 CDNA.
                                                                                                                     ADF94595 standard; CDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 CDNA.
220030964-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA.
US2003207356-Al.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003194778-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADG60928 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA,
US2003207390-A1.
                                                                                                                                                                                                                                                  ADG07217 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003207350-Al.
                                                                                                                                                                                                                                                                                                                                DB 12;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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US2003207358-Al.
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Human PRO polynucleotide #65.
US2003096966-A1.
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06-NOV-2003.
(GETH ) GENENTECH INC.
100.0%;
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22-MAY-2003.
(GETH ) GENENTECH INC.
MATCh larity 100.0%;
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06-NOV-2003.
(GETH ) GENENTECH INC.
MATCh '12-ity 100.0%;
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06-NOV-ZOGO,
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(GETH ) GENENTECH INC.
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22 MAY-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Best Local Similarity
RESULT 389
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Best Local Similarity
RESULT 382
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RESULT 365
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ID ADG609
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Score 2458; DB 12; Length 2458;
Pred. No. 0;
Length 2458;
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Novel human secreted and transmembrane protein PRO363 CDNA.
                                  AUGS6920 standard; CDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 CDNA.
                                                                                                                                                                                          ADG55816 standard, cDNA; 2458 BP. Novel human secreted and transmembrane protein PRO363 cDNA. US2003207365-Al.
                                                                                                                                                                                                                                                                                                                                                 ADGS8576 standard; CDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 CDNA.
US2003207368-A1.
06-NOV-2003.
(GETH) GENENTECH INC.
100.0%; Score 2458; DB 12; Length 3
st Local Similarity 100.0%; Pred. No. 0;
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Novel human secreted and transmembrane protein PRO363 cDNA; 1 US2003207420-A1.

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1 (GETH ) GENENTECH INC.

Query Match

100.0%; Score 2458; DB 12; Length 2
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ID ADG58024 standard; CDNA; 2458 BP.
DE Novel human secreted and transmembrane protein PRO363 CDNA
PN US2003207363-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003096965-Al.
Score 2458; DB 12;
Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Length 2458;

Length 2458;

Length 2458;

DB 12;

Best Loca RESULT 399

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AUG56368 standard; CDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 CDNA, 02203220736-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA.
US2003207367-A1.
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Human CDNA encoding secreted/transmembrane protein, PRO363
2020216560-Al.
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Novel human secreted and transmembrane protein PRO363 CDNA.
US2003207429-A1.
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Human cDNA encoding secreted/transmembrane protein PRO363
US2003224358-A1.
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                                   DB 12;
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GENENTECH INC.
100.0%; Score 2458;
Adminarity 100.0%; Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Human PRO polynucleotide #194.
US2003194793-A1.
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Human PRO polynucleotide #194.
US2003022331-A1.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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RESULT 409
                                   Query Match
Best Local Similarity
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   20-NOV-2003
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                                                                  DB 12; Length 2458;
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Human cDNA encoding secreted/transmembrane protein, PRO363.
US2003207803-A1.
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Human cDNA encoding secreted/transmembrane protein, PRO363.
US2003215908-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA.
US2003207419-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003207414-A1.
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Human cDNA encoding secreted/transmembrane protein, PRO363.
US2003216305-A1.
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Muman cDNA encoding secreted/transmembrane protein, PRO363
US2003215905-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003207416-A1.
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Human cDNA encoding secreted/transmembrane protein PRO363
US2003228656-A1.
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(GETH) GENENTECH INC.
ery Match 100.0%; Score 2458;
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Pred. No. 0;
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Human PRO polynucleotide #194.
US2003077723-A1.
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) GENENTECH INC.
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11-DEC-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.

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(GETH ) GENENTECH INC.
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RESULT 407
                                                                  Query Match
Best Local Similarity
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Best Local Similarity
RESULT 406
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Best Local S
RESULT 400
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Length 2458;

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Length 2458;

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Best Local Similarity 100.0%; Pred. No. 0; RESULT 426
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A (GETH ) GENENTECH INC.
Query Match 100.0%; Score 2458; DB 12; Length 2458;
     Score 2458; DB 12; Length 2458;
Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 12; Length 2458;
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                                                                                                                                                                                   ADG50855 standard; cDNA; 2458 BP.

Human cDNA encoding secreted/transmembrane protein, PRO363.
US2004005312-A1.
08-JAN-2004.
(GETH ) GENENTECH INC.
100.0%; Score 2458; DB 12; Length; St Local Similarity 100.0%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADG58799 standard; cDNA; 2458 BP.

Human cDNA encoding secreted/transmembrane protein, PRO363
B2004005657-A1.

08-JAN-2004.

(GETH ) GENENTECH INC.

100.0%; Score 2458; DB 12; Length Set Local Similarity 100.0%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADG34125 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2004006206-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human cDNA encoding secreted/transmembrane protein, PRO363 US2004006219-A1.
                                                   ADG59752 standard; cDNA; 2458 BP.

Novel human secreted and transmembrane protein PRO363 cDNA US2003207369-A1.
06-NOV-2003.
                                                                                                                                      DB 12;
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Pred. No. 0;
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Pred. No. 0;
                                                                                                                                      Score 2458;
Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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Pred. No. 0;
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US2003207361-A1.
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22-MBV. ...
                                                                                                                                                                                                                                                                                                                       ADH43496 standard; cDNA; 2458 BP.
Human PRO polynucleotide #32.
US2003224984-A1.
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                                                                                    US2003203
06-NOV-2003.
(GETH ) GENENTECH INC.
100.0%; S
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22-MAY-2003.
(GETH) GENENTECH INC.
100.0%;
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08-JAN-2004.
(GETH ) GENENTECH INC.
Match '---tev 100.0%;
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06-NOV-2003.
(GETH ) GENENTECH INC.
Match '174fty 100.0%;
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Human PRO polynucleotide #65.
US2004019183-A1.
     100.0%;
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04-DEC-2003.
(GETH ) GENENTECH INC.
--- Match ''--itv 100.0%;
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(GETH ) GENENTECH INC.
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2004.

AH ) GENENTECH LATAICH LOCAL SIMILARITY

RESULT 421

ID ADG34125 Stand*

PN US20040*

PD , 08-
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Best Local Similarity
RESULT 423
ID ADIB1176 standard; cl
CDNA encoding human
PN US2003207361-A1.
PD 06-NOV-2003.
PA (GETH ) GENENTECH IN
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RESULT 419
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RESULT 422
                 Best Local Similarity RESULT 417
                                                                                                                                  Query Match
Best Local Similarity
RESULT 418
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Best Local Si
RESULT 420
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      Query Match
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ADM25280 standard; cDNA; 2458 BP.
Human neurotrimin homologue related nucleotide sequence SEQ ID NO:58.
EP1386931-A1.
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Novel human secreted and transmembrane protein PRO363 CDNA.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003207383-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA
US2003096961-A1.
                                                                                                                                                  ADG09919 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2004009548-A1.
15-JAN-2004.
(GETH) GENENTECH INC.
17-JAN-2004.
18-JAN-2004.
100.0%; Score 2458; DB 12; Length;
18-Local Similarity 100.0%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                       ADII5390 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA US2003207382-A1.
06-NOV-2003.
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Novel human secreted and transmembrane protein PRO363 cDNA
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Novel human secreted and transmembrane protein PRO363 cDNA
US2004044179-A1.
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Novel human secreted and transmembrane protein PRO363 cDNA
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Pred. No. 0;
                                                  GETH) GENENTECH INC.
(GETH) GENENTECH INC.
(ery Match, "Amilarity 100.0%; Pred. No. 0;
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Pred. No. 0;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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Query Match
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Best Local Similarity
                                                                                                             Best Local Similarity RESULT 427
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RESULT 428
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                                                                                               Query Match
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AD195819 standard; cDNA; 2458 BP.
cDNA encoding human PRO polypeptide #194.
US2003077659-A1.
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                                                                                                            DB 12; Length 2458;
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US2004063921-A1.
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(GETH.) GENENTECH INC.

ry Match

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roman Similarity 100.0%; Pred. No. 0;
               ADJ77493 standard; cDNA; 2458 BP.

Human PRO polymucleotide #194.
26-FEB-2004.
(GETH ) GENENTECH INC.
100.0%; Score 2458;
ELocal Similarity 100.0%; Pred. No. 0;
                                                                                                                                                                                                                                    (GETH ) GENENTECH INC.
ry Match 100.0%; Score 2458;
t Local Similarity 100.0%; Pred. No. 0;
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ry Match
100.0%; Score 2458;
t Local Similarity 100.0%; Pred. No. 0;
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(GETH ) GENENTECH INC.
100.0%; Score 2458;
st Local Similarity 100.0%; Pred. No. 0;
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Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADJ65615 standard; cDNA; 2458 BP.
cDNA encoding human PRO polypeptide #194.
US2004038335-A1.
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CDNA encoding human PRO polypeptide #194.
US2004048333-A1.
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cDNA encoding human PRO polypeptide #194.
US2004058424-A1.
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cDNA encoding human PRO polypeptide #194.
US2004077064-A1.
                                                                                                                                                            ADK82841 standard; cDNA; 2458 BP. Human PRO polynucleotide #32. US2004043927-A1. 04-MAR-2004.
                                                                                                                                                                                                                                                                                                              ADK66605 standard, cDNA, 2458 BP.
Human PRO polynucleotide #65.
US2004044180-A1.
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22-APR-2004.
(GETH ) GENENTECH INC.
100.0%;
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(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
                                                                                                                      Best Local Similarity RESULT 436
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RESULT 442
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Best Local Similarity
RESULT 444
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RESULT 435
1D ADJ774
DE Human
PN US2004
PD 26-FEB
PA (GETH
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(HOFF ) HOFFMANN LA ROCHE & CO AG F.
ry Match
199.6%; Score 2448.4; DB 12; Length 5120;
t Local Similarity 99.8%; Pred. No. 0;
                                                   Length 2458;
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Pred. No. 0;
                                                                                                   AD196371 standard; cDNA; 2458 BP.
Novel human secreted and transmembrane protein PRO363 cDNA
US2003207354-A1.
                                                                                                                                                                                                                                                           Human soft tissue sarcoma-upregulated DNA - SEQ ID 7540 WO2004048938-A2.
                                                     DB 13;
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                                                                                                                                                                                           DB 13;
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CDNA encoding human PRO protein, Seq ID No 129.
WO200208288-A2.
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Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            73.4%; Score 1804.8; 99.7%; Pred. No. 0;
                                                     Score 2458;
Pred. No. 0;
                                                                                                                                                                                       100.0%; Score 2458;
100.0%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred. No. 0;
                                                                                                                                                                                                                                                                                                                               99.7%; Score 2450; 99.8%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 1848;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human A236 polypeptide encoding cDNA. WO200069885-A2.
                                                                                                                                                                                                                                                                                                                                                                                 ADO36719 standard; DNA; 5120 BP. .
Human UKW DNA sequence, SEQ ID NO:11.
EP1416279-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antipsoriatic cDNA sequence #909. 08-APP-^^-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABS70504 standard; cDNA; 2184 BP.
Human bone remodelling gene #161.
US6426186-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAD44196 standard; cDNA; 1948 BP
                                                                                                                                                                                                                                         ADQ24720 standard; DNA; 3120 BP.
                                                                                                                                                                                                                                                                                             10-JUN-2004.
(PROT-) PROTEIN DESIGN LABS INC.
US20030,...
24-APR-2003.
(GETH ) GENENTECH INC.
100.0%;
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Best-Local Similarity 100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (INCY-) INCYTE GENOMICS INC. ry Match 80.8%; t Local Similarity 99:98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      75.2%;
                                                                                                                                                         06-NOV-2003.
(GETH ) GENENTECH INC.
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(GETH ) GENENTECH INC.
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                                                 Ouery Match
Best Local Similarity
RESULT 445
                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                           __Best_Local Similarity_RESULT 447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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RESULT 452
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(LEIB/)
(BOSS/)
(PANY/)
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Query Match
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Human musculoskeletal system related polynucleotide SEQ ID NO 3153.
#O200155367-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADJ30526 standard; DNA; 5183 BP.
Human musculoskeletal system-associated genomic DNA - SEQ ID 3153.
US2004009488-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 1942;
                                                                                                   DB 6; Length 1948;
                                                                                                                                                                                                                                                                                                                                                                                                                DB 4; Length 1948;
                                                                                                                                                                                                                                                         DB 4; Length 1948;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABX59776 standard; cDNA; 5183 BP.
cDNA encoding novel human musculoskeletal system antigen #2120.
US2002147140-A1.
10-OCT-2002.
(ROBE/) ROSEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 4; Length 1948
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 5183;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-NOV-2000.
(MILL-) MILLENNIUM PHARM INC.
ery Match 45.6%; Score 1122; DB 4; Length 1122;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 4; Length 5183;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human soft tissue sarcoma-upregulated DNA - SEQ ID 6121
WO2004048938-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-JUN-2004.
(PROT-) PROTEIN DESIGN LABS INC.
ST.9%; Score 1422.4;
lery Match
57.9%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                 MCC-2000.
(MILL-) MILLENNIUM PHARM INC.
(MILL-) MILLENNIUM PHARM INC.
73.4%; Score 1803.2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MILL-) MILLENNIUM PHARM INC.

(MILL-) MILLENNIUM PHARM INC.

73.4%; Score 1803.2;

ery Match 79.6%; Pred. No. 0;
                                                                                                 73.4%; Score 1804.8; 99.7%; Pred. No. 0;
                                                                                                                                                                                                                                                       73.4%; Score 1803.2; 99.6%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JAN-2004.
(HUMA-) HUMAN GENOME SCI INC.
SHY MACCh 58.6%; Score 1441;
STY MACCh 99.7%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PD 02-AUG-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 58.6%; Score 1441;
Best Local Similarity 99.7%; Pred. No. 0;
RESULT 457
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             58.6%; Score 1441; 99.7%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAC84384 standard; cDNA; 1122 BP.
Human A236 polypeptide coding sequence.
WO200069885-A2.
                                                                                                                                          Luan standard; CDNA; 1948 BP. Human A236 variant 3 CDNA. 23-NOV-^^^
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAC84414 standard; cDNA; 1948 BP. Human A236 variant 2 cDNA. WO200069885-A2.
                                                                                                                                                                                                                                                                                                              AAC84413 standard; cDNA; 1948 BP.
Human A236 variant 1 cDNA.
WO200069885-A2.
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                                                                                                                                                                                                                 (MILL-) MILLENNIUM PHARM INC.
(BARN/) BARNES I M.
(FRAS/) FRASER C C.
(WRIG/) WRIGHTON N.
(MYER/) MYERS P S.
(KING/) KINGSBURY G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
RESULT 456
ID AAL36788 standard; DI
DE Human musculoskeletal
PD WO200155367-A1.
PD 02-AUG-2001.
PA (HUMA-) HUMAN GENOME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
RESULT 459
ID 'ADQ23301 standard; DN
DE Human soft tissue sar
PN WO2004048938-A2.
PD 10-UUN-2004.
PA (PROT-) PROTEIN DESIG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
RESULT 458
ID ADJ30526 standard; DI
DE Human musculoskeleta
PN US2004009488-A1.
PD 15-JAN-2004.
PA (HUMA-) HUMAN GENOME
                                                                                               Query Match
Best Local Similarity
RESULT 453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ouery Match
Best Local Similarity
RESULT 460
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Best Local Similarity
RESULT 454
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Best Local Similarity
RESULT 455
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Length 1949;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 1949;
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27-SEP-2001.
(MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
Query Match 23.4%; Score 575.2; DB 5; Length 1233;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 1122;
                                                                                              Length 1949;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 1949;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 42.4%; Score 1043.4; DB 4; 79.2%; Pred. No. 4.7e-276;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         42.4%; Score 1043.4; DB 4; 79.2%; Pred. No. 4.7e-276;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               42.4%; Score 1043.4; DB 4; 79.2%; Pred. No. 4.7e-276;
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AAC84391 standard; cDNA; 1949 BP.
Mouse A236 polypeptide encoding cDNA.
W0200069885-A2.
23-NOV-2000.
(MILL-) MILLENNIUM PHARM INC.
ETY MATCH
TY MATCH
TY NOV-2000 TO 38; Pred. NO. 1.7e-276;
                                                                                                                                                                                                                                                                                                                                                                                                                                     42.5%; Score 1045; DB 6; 79.3%; Pred. No. 1.7e-276;
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WO200166720-Al.
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Human ovarian cancer DNA marker #20546.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAC84417 standard; cDNA; 1949 BP.
Mouse A236 variant 2 cDNA.
WO200069885-A2.
                                                                                                                                                        AAD44197 standard; cDNA; 1949 BP.
Mouse A236 cDNA.
US2002055139-A1.
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ID AAC84416 standard; cDNA; 1949 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAC84418 standard; cDNA; 1949 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-NOV-2000.
(MILL-) MILLENNIUM PHARM INC.
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(MILL-) MILLENNIUM PHARM INC.
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Query Match
Best Local Similarity 79.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse A236 variant 3 cDNA. WO200069885-A2.
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WO200069885-A2.
23-NOV-2000.
                                                                                                                                                                                                                                 (HOLT/) HOLTZMAN D A (SHAK/) SHARP J D. (LEIS/) LEIBY R R. (BOSS) BOSSONE S. (PANY/) PAN Y. (BARN/) BARNES T M. (RRAS/) FRASER C C. (WRIG/) WRIGHYON N. (WYER/) MYERS P S.
                                                                                                                                                                                                                                                                                                                                                                                          (MYER/) MYERS P S.
(KING/) KINGSBURY G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
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Best Local Similarity
RESULT 463
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(TSUR/) TSURUGA H.
                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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Length 319;

Query Match

us-10-002-967a-58.rng.spdi

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ADJ08178 standard; DNA; 319 BP.
Human cardiovascular system associated polypeptide-related DNA SeqID1566.
US2004005575-A1.
                                                                                                                                  Score 79.4; DB 11; Length 43454; Pred. No. 6.1e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-SEP-2002.
(ORTH ) ORTHO CLINICAL DIAGNOSTICS INC.
ery Match 2.9%; Score 71.6; DB 6; Length 136328;
                                                                                                                                                                                           AAS36066 standard; DNA; 319 BP.
Human cardiovascular system antigen genomic DNA SEQ ID No 1566.
WO20015531-A2.
02-ANG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAL46723 standard; DNA; 1000 BP.
Human myosin-like gene hGDMLP-1 transcriptional control region.
EP1158049-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-JAN-2004.
(HUMA-) HUMAN GENOME SCI INC.
2.9%; Score 72.2; DB 13; Length 319;
ery Match
2.9%; Pred. No. 3.8e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.9%; Score 72.2; DB 10; Length 319; 78.9%; Pred. No. 3.8e-09;
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Pred. No. 2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human nervous system related polynucleotide SEQ ID NO 9581 WO200159063-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABZ35015 standard, cDNA, 136328 BP.
Human gene expression profile polynucleotide SEQ ID NO 127
WO200274979-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human GDMLP-1 putative promoter sequence SEQ ID NO:15752 WO200192524-A2.
                                                                                                                                                                                                                                                                                                                                                               ADE46760 standard; DNA; 319 BP.
Human cardiovascular system related genomic DNA #326.
US2003059908-A1.
                                                                                                                                                                                                                                                                                                     2.9%; Score 72.2; DB 4; 78.9%; Pred. No. 3.8e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.9%; Score 71.6; DB 5; 70.9%; Pred. No. 9.5e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    / Match 2.9%; Score 72; DB 11;
Local Similarity 93.8%; Pred. No. 4.6e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 70.6; DB
Pred. No. 2e-08;
                        ACN43906 standard; DNA; 43454 BP. Human genomic sequence hCG15249. 402003073826-A2. 12-SEP-2003. (SAGR-) SAGRES DISCOVERY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GENO-) GENOME THERAPEUTICS CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADP75185 standard; DNA; 22693 BP.
Human NRG2 gene fragment #1.
WO2003031594-A2.
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(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                               (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                               27-MAR-2003.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                             3.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABA17250 standard; DNA; 839
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-DEC-2001.
(AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity
RESULT 484
                                                                                                                                               Best Local Similarity RESULT 479
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Best Local Similarity
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Best Local Similarity
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(AEOM-) AEOMICA INC.
                                                                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABN15760 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 487
                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                vuery Match 14.3%; Score 352.6; DB 11; Length 1496; Best Local Similarity 98.9%; Pred. No. 5.9e-86; RESULT 473
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8.3%; Score 203.2; DB 5; Length 1602; 98.2%; Pred. No. 8.1e-45;
                                                                                                                                                                                                                                                                                                                         Length 425;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 405;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 295;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 295;
                                                                                                                                                                                                                                                                                                                                                                                                                                                O 10-JUN-2004.

A (PROT-) PROTEIN DESIGN LABS INC.

Query Match 15.5%; Score 380; DB 12; Length 392;

Best Local Similarity 98.7%; Pred. No. 8e-94;
                                                                                                                                                 17.3%; Score 426; DB 8; Length 426; 100.0%; Pred. No. 1.8e-106;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 332
                                                                                                                                                                                                                                                                                                                                                                                                  Human soft tissue sarcoma-upregulated DNA - SEQ ID 3569 WO2004048938-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human soft tissue sarcoma-upregulated DNA - SEQ ID 1750 WO2004048938-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-SEP-2001.
(MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
ery Match 13.8%; Score 339.4; DB 5;
ery incal Similarity 95.6%; Pred. No. 1.2e-82;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-SEP-2001.
(MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
217 Match 9.0%; Score 221.6; DB 5;
21 Tocal Similarity 91.4%; Pred. No. 2.7e-50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PD 27-SEP-2001.

PA (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.

Query Match 9.0%; Score 221.6; DB 5;

Best Local Similarity 91.4%; Pred. No. 2.7e-50;

RESULT 477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cDNA; 1602 BP. human diagnostic protein #8723
                                                                                                                                                                                                                                                                                                                     16.7%; Score 411.4; DB 8; 99.5%; Pred. No. 1.9e-102;
    98.4%; Pred. No. 2.8e-147;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLOUS TO TO TO THE TOTAL THE T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACN90389 standard; DNA; 1496 BP.
Breast cancer related marker, seq id 11539.
US2003099974-A1.
                          CDNA encoding novel human polypeptide #70.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADL41533 standard; DNA; 405 BP.
Human ovarian cancer DNA marker #15423.
WO200170979-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADI76297 standard; DNA; 295 BP.
Human ovarian cancer DNA marker #9039.
WO200170979-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADI69964 standard; DNA; 295 BP.
Human ovarian cancer DNA marker #2706
WO200170979-A2.
                                                                                                                                                                                                             ACD05879 standard; cDNA; 425 BP.
Novel human contig #53.
WO2003023013-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADQ18931 standard; DNA; 332 BP.
                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-MAY-2003. (MILL-) MILLENNIUM PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                     DNA; 392
Best Local Similarity RESULT 469
                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity RESULT 474
                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
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Best Local Similarity
RESULT 475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA encoding novel WO200175067-A2.
                                                                                                     20-MAR-2003.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                        20-MAR-2003.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                     ADQ20749 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAS72919 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
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                                                                                                                                               Query Match
Best Local Si
RESULT 470
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Query Match

Length 22693;

Length 839

Length 1000

9

Query Match

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(AVAL-) AVALON PHARM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18-JUL-2002
                           13-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                 02-AUG-2001
                                                                                                                                                              13-DEC-2001
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                                                           Query Match
                                                                                            RESULT 496
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PA PA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               . AAK67165 standard; DNA; 6782 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:21977.
WO200157182-A2.
Human GDMLP-1 transcriptional control region cDNA SEQ ID NO:15752. US2004137589-A1. 15-JUL-2004. (GTYY) GU Y. (JTYY) JI Y. (TTY) JI Y. (HENN) PENN S G. (HANZ) HANZEL D K.
                                                                                                                                                                                                                                                                                                                                                                                                                Human immune/haematopoietic antigen encoding cDNA SEQ ID NO:1065 WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 70.4; DB 10; Length 96596;
Pred. No. 2.8e-07;
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                                                                                                                                                                                                               / Match 2.9%; Score 70.6; DB 13; Length 1000; Local Similarity 74.5%; Pred. No. 2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 6782;
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Colon adenocarcinoma related gene sequence SEQ ID NO:284
WO200194629-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADA02504 standard; DNA; 96596 BP.

Human BACH2 carcinoma associated gene, SEQ ID NO:1022.
WO2003057146-A2.

17-UTL-2013.

(SAGR-) SAGRES DISCOVERY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABD33066 standard; DNA; 87847 BP.

Human cancer-associated genomic DNA HD23-013.

W02004074320-A2.

W02SEP-2004.

(SAGR-) SAGRES DISCOVERY INC.

2.9%; Score 70.4; DB 13;

st Local Similarity 92.5%; Pred. No. 2.7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 70.4; DB 10;
Pred. No. 2.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Weace-2001.

09-AUG-2001.

(HUMA-) HUMAN GENOME SCI INC.

2.9%; Score 70.4; DB 4;

ery Match

2.9%; Pred. No. 6.5e-08;
                                                                                                                                                                                                                                                                                                                                                   2.9%; Score 70.6; DB 6; 74.5%; Pred. No. 2.9e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                           09-ACC-2001.
(HUMA-) HUMAN GENOME SCI INC.
(HUMA-) HUMAN GENOME SCI INC.
(ery Match 2.9%; Score 70.4; DB 4;
                                                                                                                                                                                                                                                       Human GDMLP-1 putative promoter sequence. WO200192524-A2. 65-DEC-2001. EDMC-2001. EDMC-2001. Y.Y. March
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADE95752 standard; DNA; 96596 BP.
Human BACH2 gene genomic DNA sequence.
WO2003039484-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA; 96596 BP
                                                                                                                                                                                                                                                                                                                                                                                                   CDNA; 758 BP
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92.5%;
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Local Similarity 92.5%;
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N WC2003008583-A2.
D 30-JAN-2003.
A (SAGR-) SAGRES DISCOVERY.
Query Match 2.9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-MAY-2003.
(SAGR-) SAGRES DISCOVERY.
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RESULT 490
ID AAK67165 standard; DN
DE Human immune/haematop
PN WO200157182-A2.
PD 09-AUG-2001.
PA (HUMA-) HUMAN GENOME
                                                                                                                                                                                               SHANNON M E.
                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
RESULT 489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity RESULT 491
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                   AAK56005 standard;
                                                                                                                                                              RANK D.
CHEN W.
                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                                                                                                Query Match
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                                                                                                                                                            (RANK/) I
(CHEN/) (SHAN/)
                                                                                                                                                                                                                                              RESULT 488
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ADJ08179 standard; DNA; 299 BP.
Human cardiovascular system associated polypeptide-related DNA SeqID1567.
US2004005575-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADB82573 standard; cDNA; 659 BP.
Human cDNA sequence useful for the treatment of cancer (SeqID 885).
Length 149480;
                                                                                                                                                                  Length 149480;
                                                                                                                                                                                                                                                                                                                                  Length 149480;
                                                                                                                                                                                                                                                                                                                                                                                   AAS36067 standard; DNA; 299 BP.
Human cardiovascular system antigen genomic DNA SEQ ID No 1567
WO200155321-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.9%; Score 70.2; DB 13; Length 299. 78.5%; Pred. No. 1.3e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 34641;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 299;
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                                                                                                                                                                                                                         ABL61948 standard, DNA, 149480 BP.
Colon adenocarcinoma related gene sequence SEQ ID NO:285.
W020019429-A2.
13-DEC-2001.
(AVAL-) AVALON PHARM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADE46761 standard; DNA; 299 BP.
Human cardiovascular system related genomic DNA #327
US2003059908-A1.
                                                             ABL68365 standard; DNA; 149480 BP.
Kidney cancer related gene sequence SEQ ID NO:6702
WO200194629-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABQ89631 standard; cDNA; 730 BP.
Human prostate expressed polynucleotide SEQ ID NO
WQ200255700-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-MAR-2003.
(HUMA-) HUMAN GENOME SCI INC.
2.9%; Score 70.2; DB 10;
ery Match 2.9%; Pred. No. 1.3e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lung cancer related gene sequence SEQ ID NO:4137.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lung cancer related gene sequence SEQ ID NO:4814 WO200194629-A2.
    2.9%; Score 70.4; DB 6; 92.5%; Pred. No. 3.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.9%; Score 70.2; DB 4; 78.5%; Pred. No. 1.3e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 6; I
2.1e-07;
                                                                                                                                                                  Score 70.4; DB 6;
Pred. No. 3.6e-07;
                                                                                                                                                                                                                                                                                                                              Query Match 2.9%; Score 70.4; DB 6; Best Local Similarity 92.5%; Pred. No. 3.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.8%; Score 70; DB 9;
84.0%; Pred. No. 2.3e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.8%; Score 70; DB 6;
84.0%; Pred. No. 2.4e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 70;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABL65800 standard; DNA; 34641 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABL66477 standard; DNA; 34641 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-JAN-2004.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                  2.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (AVAL-) AVALON PHARM.
                                                                                                                                                 (AVAL-) AVALON PHARM.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CHIR ) CHIRON CORP. (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CHIR ) CHIRON CORP. (HYSE-) HYSEQ INC.
                  Best Local Similarity
                                                                                                                                                                                     Best Local Similarity RESULT 497
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 501
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
RESULT 500
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Length 4666;

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Query Match
Best Local Similarity
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Best Local Similarity
              Best Local Similarity RESULT 512
                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 520
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-JAN-2003
                                                                                                     09-AUG-200
                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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                                                                                                                                       Query Match
   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                         RESULT 513
                                                                                                                                                                                                                                                                                                              RESULT 514
                                                                                                                                                                                                                                                                                                                                                                                            ADJ12490 standard; DNA; 45698 BP.
DNA fragment of a BAC clone that encodes a human secreted protein Seq344.
                                                                                                                                              12-SEP-2003.
(SAGR-) SAGRES DISCOVERY.
ery Match
ery Match
-- Tonal Similarity 91.4%; Pred. No. 7.46-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.8%; Score 69.4; DB 11; Length 196686; 92.4%; Pred. No. 7.9e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 228139;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.8%; Score 69.4; DB 12; Length 45698; 92.4%; Pred. No. 3.5e-07;
                                                                                                                                                                                                                                                                                                                                              Length 933;
                                                   2.8%; Score 70; DB 6; Length 34641; 71.1%; Pred. No. 2.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 4666;
                                                                                                                                                                                                                                     AD041322 standard; cDNA; 933 BP.
Human cDNA probe useful for disease diagnosis #473.
WO2004046382-A2.
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(12-SEP-2003.

(SAGR-) SAGRES DISCOVERY.

2.8%; Score 69.4; DB 11;

2.8%; Score 69.4; DB 11;

2.8%; Pred. No. 8.6e-07;
                                                                                                                                                                                                                                                                                                                                            2.8%; Score 69.4; DB 12; 92.4%; Pred. No. 4.1e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.8%; Score 69; DB 7; 183.9%; Pred. No. 1.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADP44060 standard; cDNA; 4666 BP.
Human CD11C encoding cDNA SEQ ID NO:13.
WO2004053094-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADS99918 standard; cDNA; 4666 BP.
Human integrin, alpha X (ITGAX), cDNA.
US2003148327-A1.
                                                                                                     DNA; 276276 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ACN44002 standard; DNA; 228139 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACN44170 standard; DNA; 196686 BP.
                                                                                                                 Human genomic sequence hCG17121.
WO2003073826-A2.
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WO2003073826-A2.
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WO2003073826-A2.
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(SAGR-) SAGRES DISCOVERY.
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(OLEK/) OLEK A.
(PIEP/) PIEPENBROCK C.
(BERL/) BERLIN K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FLORENCE K A. GREENE J M. YOUNG P E. PERRIE A M.
                 13-DEC-2001.
(AVAL-) AVALON PHARM.
                                                                                                                                                                                                                                                                                                         (DIAG-) DIAGENIC AS. (JONE/) JONES E L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BREWER L A.
DUAN R D.
RUBEN S M.
                                                                 Local Similarity
                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       YU G.
FLORENCE C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                 ACN44350 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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(PPDP-) PPD DEV LP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                ROSEN C A
                                                                                                                                                                                                                                                                                                                                                                                                                                  JS2004010132-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EBNER R.
OLSEN H.
WO200194629-A2.
                                                                                                                                                                                                                                                                                              03-JUN-2004
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(ROSE/) ROSI
                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (EBNE/)
(OLSE/)
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RUBE/)
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ABK42318 standard; DNA; 251 BP.
Genomic sequence #217 encoding novel human connective tissue polypeptide.
WO200155343-A1.
                                                                AAK65953 standard; DNA; 158 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:20765.
WO200157182-A2.
                                                                                                                                                                                                                                                      AAK65950 standard; DNA; 158 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:20762.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                              ABZ74262 standard; DNA; 232 BP.
Secreted protein gene 215 genomic fragment HNGAM58, SEQ ID NO:1409.
WO200277013-A2.
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Human colon cancer related nucleotide sequence SEQ ID NO:325.
WQ20<u>0</u>229086-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.8%; Score 68.8; DB 10; Length 232 91.2%; Pred. No. 2.8e-08;
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                                                                                                                                                                                      Length 158
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WO200277186-A2.
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Human pancreatic cancer expressed cDNA SEQ ID NO 2882
WO200260317-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human secreted protein-related DNA sequence #384, W02003004623-A2.
                                                                                                                                                                                    2.8%; Score 68.8; DB 4;
91.2%; Pred. No. 2.2e-08;
                                                                                                                                                                                                                                                                                                                      09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.8%; Score 68.8; DB 4;
ery Match 91.2%; Pred. No. 2.2e-08;
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02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
(HOMA-) HUMAN GENOME SCI INC.
2.8%; Score 68.8; DB 4;
ery Match
2.9%; Pred. No. 2.9e-08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.8%; Score 68.8; DB 8; 91.2%; Pred. No. 2.8e-08;
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2.8%; Score 69; DB 12;
83.9%; Pred. No. 1.3e-07;
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Connective tissue related genomic DNA #217.
US2003054375-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABZ67828 standard; DNA; 232 BP.
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(HUMA-) HUMAN GENOME SCI INC.
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(HUMA-) HUMAN GENOME SCI INC.
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(HUMA-) HUMAN GENOME SCI INC.
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CORI-) CORIXA CORP.
Query Match
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Best Local Similarity 91.2%; Pred. No. 1.6e-07; RESULT 530
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA; 7720 BP.
                                                                                                                                                                                             DNA; 6902 BP
                                                                                                                                                                                                                                                02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
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Best Local Similarity 91.2%;
RESULT 536
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(AVAL-) AVALON PHARM.
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(AVAL-) AVALON PHARM.
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(AVAL-) AVALON PHARM.
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Best Local Similarity
                                                                                                                                                                                                                                                                                               Best Local Similarity RESULT 532
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                                                                                                                                                                                         ABK42132 standard;
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:29911.
909-AUG-2001.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:31441.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                Length 1406;
                                                                                                                                         Length 1001;
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                                                                                                                                                                                                                                                                                     Length 1406;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADS3253 standard; DNA; 2723 BP.
ADS3253 standard; DNA; 2723 BP.
WO2004082463-A2.
30-SEP-2004.
(MEDI-) MEDICAL COLLEGE OHIO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADS32763 standard, DNA; 2723 BP.
Human FR-alpha P1 and P4 promoter gene fragment SEQ ID NO:16.
WO2004082463-A2.
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                                       ADO34635 standard; DNA; 1001 BP.
Human SLIT-3 gene microsatellite polymorphic region #200.
WO2004042358-A2.
       yene microsatellite polymorphic reg. 21-MAY-2004.

PA (DECO-) DECODE GENETICS EHF.
Query Match
PACT8806 standard; 91.2%; Pred. No. 6.2e-08; RESULT 5.2

ID ADC78806 standard; DNA; 1406 BP.
PD Human PRO protein coding serv.
PN WO20034984-A2.
PA (CELL)
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(MEDI-) MEDICAL COLLEGE OHIO.
(REDI-) MATCh 2.8%; Score 68.8; DB 13;
ery Match 91.2%; Pred. No. 1.1e-07;
                                                                                                                                                                                                                                                                                  2.8%; Score 68.8; DB 10; 91.2%; Pred. No. 7.5e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                           Match 2.8%; Score 68.8; DB 10; Local Similarity 91.2%; Pred. No. 7.5e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Match 2.8%; Score 68.8; DB 10; Local Similarity 91.2%; Pred. No. 9.3e-08;
                                                                                                                                         DB 12;
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91.2%; Pred. No. 1.1e-07;
Best Local Similarity 91.2%; Pred. No. 3.3e-08. RESULT 521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADB63519 standard; cDNA; 2091 BP.
Human cDNA encoding clone TESTI20254090.
EP1308459-A2.
                                                                                                                                                                                                                                                                                                                                                         Human PRO protein coding sequence #61.
WO2003034984-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADF48653 standard; cDNA; 2820 BP.
Human PP7518 cDNA #SEQ ID 10.
CN1369505-A.
                                                                                                                                                                                                                                                                                                                                        ADC78892 standard; DNA; 1406 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-SEP-2002.
(SHAN-) SHANGHAI INST ONCOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HUMA-) HUMAN GENOME SCI INC.
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                           Best Local Similarity
RESULT 523
ID ADC78892 standard; DN
DE Human PRO protein cod
PN WO2003034984-A2.
PD 01-MAY-2003.
PA (GETH ) GENENTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 526
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Best Local Similarity
RESULT 527
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RESULT 5
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ID AD
DE Hu
PN WO
PD 30
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Genomic sequence #31 encoding novel human connective tissue polypeptide.
WO200155343-A1.
                                                                                                                             Ouery Match 2.8%; Score 68.8; DB 12; Length 5638;
Best Local Similarity 91.2%; Pred. No. 1.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.8%; Score 68.8; DB 6; Length 7720; 91.2%; Pred. No. 1.9e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 6902;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kidney cancer related gene sequence SEQ ID NO:6595. W0200194629-A2. 13-DEC-2001. (AVAL-) AVALON PHARM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABL68540 standard; DNA; 7720 BP.
Kidney cancer related gene sequence SEQ ID NO:6877.
WO200194629-A2.
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Kidney cancer related gene sequence SEQ ID NO:7213.
WO200194629-A2.
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Lung cancer related gene sequence SEQ ID NO:3973.
WO200194629-A2.
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WO200194629-A2.
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Best Local Similarity 91.2%; Pred. No. 1.9e-07;
RESULT 537
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(UYNE-) UNIV NEW JERSEY MEDICINE & DENTISTRY.
ery Match
st Local Similarity 91.2%; Pred. No. 1.9e-07;
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Pred. No. 1.9e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.8%; Score 68.8; DB 9; 91.2%; Pred. No. 1.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 2.8%; Score 68.8; DB 6; Best Local Similarity 91.2%; Pred. No. 1.9e-07;
                                                                                                                                                                                                                                                                                                                                     2.8%; Score 68.8; DB 4; 91.2%; Pred. No. 1.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                           ADB60288 standard; DNA; 6902 BP.
Connective tissue related genomic DNA #31.
US2003054375-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAC91210 standard; DNA; 7720 BP.
Human folate receptor I gene SEQ ID NO: 5.
WO200071754-A1.
ADD07499 standard, DNA; 5638 BP.
Human brain-derived gene SEQ ID NO:2.
24-UNN-2004.
(KAZU-) ZH KAZUSA DNA KENKYUSHO.
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Length 17239;

03-OCT-2002

Best 1

Query Match Best Local S

RESULT 542

Query Match Best Local Si RESULT 543

Query Match Best Local S

Length 17239,

Length 17239

Length 18408;

Length 18408

Length 18408

Query Match

Best Loca RESULT 548

RESULT 546
ID AAK816
DE Human
PN WO2001
PD 09-AUG
PA (HUMA-

Best Loca RESULT 547

Query Match

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Human cardiovascular system associated polypeptide-related DNA SeqID2326.
US2004005575-A1.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:20180.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.8%; Score 68.8; DB 10; Length 17239; 97.2%; Pred. No. 3e-07;
                         Human cardiovascular system antigen genomic DNA SEQ ID No 2326.
WO200155321-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.8%; Score 68.8; DB 4; Length 18564; 73.9%; Pred. No. 3.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human testicular antigen encoding DNA fragment SEQ ID NO: 2308 WO200155317-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABL97657 standard; DNA; 18408 BP.
Human testicular antigen encoding DNA fragment SEQ ID NO: 2309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AALO4753 standard; DNA; 18408 BP.
Human reproductive system related antigen DNA SEQ ID NO: 7441.
WO200155320-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAL04754 standard; DNA; 18408 BP.
Human reproductive system related antigen DNA SEQ ID NO: 7442.
WO200155320-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 18408
                                                                                                                                                                                                                                                                                                                                                  ABZ67711 standard; DNA; 17239 BP.
Human secreted protein encoding genomic DNA SEQ ID NO 1234
WO200277186-A2.
                                                                                                                                                                                                Human cardiovascular system related genomic DNA #1086 US2003059908-A1.
                                                                                                                                                                                                                                                                                       2.8%; Score 68.8; DB 10; 97.2%; Pred. No. 3e-07;
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                                                           G2-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.8%; Score 68.8; DB 4
ery Match 97.2%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.8%; Score 68.8; DB 97.2%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADJ08938 standard; DNA; 17239 BP.
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      AAS36826 standard; DNA; 17239 BP
                                                                                                                                                                           DNA; 17239 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA; 18408 BP
                                                                                                                                                                                                                                           27-MAR-2003.
(HUMA-) HUMAN GENOME SCI INC.
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(HUMA-) HUMAN GENOME SCI INC.
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                                                                                                                Query Match
Best Local Similarity
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                                                                                                                                                                         ADE47520 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT
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US2004005575-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABL55889 standard; DNA; 10012 BP.
Human small inducible cytokine subfamily A (Cys-Cys), member 24 gene.
WO200220851-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human immune/haematopoietic antigen genomic sequence SEQ ID NO:36474, WO200157182-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAK88955 standard; DNA; 12724 BP.
Human digestive system antigen genomic sequence SEQ ID NO: 2531.
WO200155314-A2.
                    AAS36827 standard; DNA; 7960 BP.
Human cardiovascular system antigen genomic DNA SEQ ID No 2327.
W0200155321-A2.
02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
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(HUMA-) HUMAN GENOME SCI INC.
ery Match
2.8%; Score 68.8; DB 4; Length 16817;
                                                                                                                                                                                                                                                                                                           Length 7960;
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                                                                                                                                Length 7960;
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Human secreted protein encoding genomic DNA SEQ ID NO 1235.
WO200277186-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAL02697 standard; DNA; 9339 BP.
Human reproductive system related antigen DNA SEQ ID NO:
WO200155320-A2.
                                                                                                                                                                                                                  Human cardiovascular system related genomic DNA #1087.
US2003059908-A1.
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ry Match

t Local Similarity 97.2%; Pred. No. 2e-07;
                                                                                                                                                                                                                                                                                                           2.8%; Score 68.8; DB 10; 97.2%; Pred. No. 2e-07;
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Best Local Similarity 91.2%; Pred. No. 2.2e-07;
RESULT 545
                                                                                                                                  2.8%; Score 68.8; DB 4;
97.2%; Pred. No. 2e-07;
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(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.8%; Score 68.8; DB 4;
ery Match 91.2%; Pred. No. 2.1e-07;
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(HUMA-) HUMAN GENOME SCI INC.
ETY MATCh 2.8%; Score 68.8; DB :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADJ08939 standard; DNA; 7960 BP.
                                                                                                                                                                                                ADE47521 standard; DNA; 7960 BP
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(GENA-) GENAISSANCE PHARM INC.
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(HUMA-) HUMAN GENOME SCI INC.
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(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                    27-MAR-2003.
A (HUMA-) HUMAN GENOME SCI INC.
Query Match
RESULT 539

ID AAS36827 standard; D
B Human cardiovascular
PN WO200155321-A2.
PD 02-AUG-2001.
PA (HUMA-) HUMAN GENOME
                                                                                                                                                                                                                                                                                                                         Local Similarity
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(HUMA-) HUMAN GENOME
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Best Local Similarity
RESULT 567
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Best Local Similarity
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Best Local Similarity
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                               WO200157182-A2.
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RESULT 572
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                                                                                                                                                                                                                                                                                                                                  Genomic sequence #32 encoding novel human connective tissue polypeptide. WO200155343-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAK81665 standard; DNA; 24908 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:36477.
W0200157182-A2.
W0200157182-A2.
(HUMA-) HUMAN GENOME SCI INC.
ery Match
2.8%; Score 68.8; DB 4; Length 24908;
st Local Similarity 91.2%; Pred. No. 3.7e-07;
          Secreted protein gene 300 genomic fragment HSQDO85, SEQ ID NO:1608
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Best Local Similarity 91.2%; Pred. No. 3.8e-07;
RESULT 563
                                                                                                                                              ADA98881 standard; DNA; 18564 BP.

Human secreted protein-related DNA sequence #474.

W02003004623-A2.

W02A04-2003.

(HUMA-) HUMAN GENOME SCI INC.

2.8%; Score 68.8; DB 8; Length 18564;

St Local Similarity 73.9%; Pred. No. 3.1e-07;
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                                                                                           Length 18564;
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Human cancer associated gene genomic sequence SEQ ID NO:58.
WO2004074321-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAL06991 standard; DNA; 32082 BP.
Human reproductive system related antigen DNA SEQ ID NO:
WO200155320-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAL51501 standard; DNA; 35057 BP.
Human pantothenate kinase 2 (PANK2) genomic DNA sequence
WQ2003008626-A2.
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PA (SAGR-) SAGRES DISCOVERY INC.
Query Match
Best Local Similarity 91.2%; Pred. No. 4.3e-07;
RESULT 565
DE Secretea process
PN W0200277013-A2.
PD 03-0CT-2002.
PD 03-0CT-2002.
PA (HUMAN GENOME SCI INC.
PA HUMAN GENOME SCI INC.
2.8%; Score 68.8; DB 8;
Corr Match 13.14 73.9%; Pred. No. 3.1e-07;
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02-AUG-2001.
(HUMAL) HUMAN GENOME SCI INC.
(HACL)
2.8%; Score 68.8; DB 4;
ery Match
2.8%; Pred. No. 3.5e-07;
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PA (HUMA-) HUMAN GENOME SCI INC.
Query Match
Best Local Similarity 91.2%; Pred. No. 3.5e-07;
RESULT 561
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02-MCJOOL.

(HUMA-) HUMAN GENOME SCI INC.

(HTMA-) HUMAN GENOME SCI INC.

2.8%; Score 68.8; DB 4;

ery Match

2.8%; Pred. No. 4.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADB60289 standard, DNA; 22161 BP.
Connective tissue related genomic DNA #32.
US2003054375-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACN44246 standard; DNA; 26047 BP. Human genomic sequence hCG20994. U2-200303926-A2. (SAGR-) SAGRES DISCOVERY.
                                                                                                                                                                                                                                                                                                                     DNA; 22161 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAK70780 standard; DNA; 37783 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UYOR-) UNIV OREGON HEALTH SCI. (REGC ) UNIV CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
RESULT 562
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity RESULT 564
                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
RESULT 559
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Best Local Similarity
RESULT 560
                                                                                                                                                                                                                                                                                                                   ABK42133 standard;
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Best Local S:
RESULT 566
ID AAK70780
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ABN89533 standard; DNA; 48436 BP.
Human corneal N-acetylglucosamine-6-sulfotransferase DNA SEQ ID NO:38.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:37440.
WO200157182-A2.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:24973.
WO200157182-A2.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:37442.
WO200157182-A2.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:36478.
WO200157182-A2.
                                                                                                                                                                                                           Human immune/haematopoietic antigen genomic sequence SEQ ID NO:31437.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                            Human immune/haematopoietic antigen genomic sequence SEQ ID NO:35725.
WO200157182-A2.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:36475.
WO200157182-A2.
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WO200157182-A2.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:25592
                                                                                                                                                                                                                                                                                                                         Length 37783;
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(HUMA-) HUMAN GENOME SCI INC.
2.8%; Score 68.8; DB 4;
ery Match 2.8%; Pred. No. 5.3e-07;
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91.2%; Pred. No. 5.3e-07;
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91.2%; Pred. No. 4.7e-07;
                                                                         SCI INC.
2.8%; Score 68.8; DB 4;
91.2%; Pred. No. 4.7e-07;
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91.2%; Pred. No. 4.7e-07;
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91.2%; Pred. No. 5.3e-07;
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                                                                                                                                                                                    AAK76625 standard; DNA; 37783 BP
                                                                                                                                                                                                                                                                                                                                                                                                           AAK80913 standard; DNA; 37783 BP.
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(HUMA-) HUMAN GENOME SCI INC.
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Best Loca RESULT 577

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2.8%; Score 68.8; DB 13; Length 106938; 91.2%; Pred. No. 8.3e-07;
                                                                                                                             2.8%; Score 68.8; DB 10; Length 68355; 91.2%; Pred. No. 6.5e-07;
                                                                                                                                                                                                                                                                              (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.

ry Match
2.8%; Score 68.8; DB 10; Length 68355;
t Local Similarity 91.2%; Pred. No. 6.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 68.8; DB 10; Length 96588;
Pred. No. 7.8e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               / Match 2.8%; Score 68.8; DB 9; Length 96588; Local Similarity 91.2%; Pred. No. 7.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human MBNL carcinoma associated gene, SEQ ID NO:1544 WO2003057146-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADM74621 standard; DNA; 96588 BP.
Human carcinoma associated (CA) nucleic acid #145.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
ry Match 2.8%; Score 68.8; DB 10;
t Local Similarity 91.2%; Pred. No. 6.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.8%; Score 68.8; DB 12; 91.2%; Pred. No. 7.3e-07;
                                                                                                                                                                                                                                                                                                                                              Human MDR1 related DNA; 68355 BP.
Human MDR1 related DNA sequence SEQ ID NO:665.
WG200313535-A2.
                                                                                                                                                                                                                 Human MDR1 related DNA sequence SEQ ID NO:665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABD33432 standard; DNA; 106938 BP.
Human cancer-associated (CA) gene HD07-081.
ID ADB87941 standard; DNA; 68355 BP.

ID ADB87941 standard; DNA; 68355 BP.

BE Human UGT1A1 gene sequence SEQ ID NO:665.

BW WO2003013536-A2.

PD 20-FEB-2003.

PD 20-FEB-2003.

PA (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CDNA; 86000 BP
                                                                                                                                                                                             DNA; 68355 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADA03026 standard; DNA; 96588 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADB72764 standard; DNA; 96588 BP
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05-JUN-2003.
(SAGR-) SAGRES DISCOVERY.
ARY MATCH 2.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.8%;
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(SAGR-) SAGRES DISCOVERY.
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(ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MORR/) MORRIS D W. (ENGE/) ENGELHARD E K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human PPAR-alpha cDNA.
US2004115637-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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593.
                                                                                                                                            Best Local Similarity RESULT 586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                  Best Local Similarity RESULT 587
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                                                                                                                                                                                             ADB96924 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADP68568 standard;
                                                                                                                                                                                                                                  WO2003013537-A2.
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WO2003008583-A2.
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                                                                                                                                                                                                                                                                                                    AAK73083 standard, DNA, 50442 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:27895.
WO200157182-A2.
                                                                                                                                                                         Human immune/haematopoietic antigen genomic sequence SEQ ID NO:42363.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADS3462 standard; DNA; 64423 BP.
Human autoimmune disease-related genomic DNA seguence - SEQ ID 1676.
WO2004083403-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human autoimmune disease-related genomic DNA sequence - SEQ ID 1668. 30-SEP-2004. APPLERA CORP.
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                                                                                        Length 48436;
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                                                                                                                                                                                                                                                             Length 50442
                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 50442
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Cancer based on CYP3A5 related polynucleotide SEQ ID NO:665.
WO2003013534-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADB20852 standard, DNA, 68355 BP. MRP1 based cancer related nucleic acid SEQ ID NO:665
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(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.8%; Score 68.8; DB 4;
ery Match 91.2%; Pred. No. 5.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 13;
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Pred. No. 6.3e-07;
                                                                                                                                                                                                        09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ETY Match 2.8%; Score 68.8; DB 4;
ery Match 91.2%; Pred. No. 5.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.8%; Score 68.8; DB 8; 91.2%; Pred. No. 5.8e-07;
                                                                                        2.8%; Score 68.8; DB 6; 91.2%; Pred. No. 5.3e-07;
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PA (SAGR-) SAGRES DISCOVERY INC.
Query Match 2.8%; Score 68.8; DB 13.
Best Local Similarity 91.2%; Pred. No. 5.9e-07;
RESULT 581
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human cancer-associated genomic DNA HD23-001.
WO2004074320-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACA60949 standard; DNA; 55827 BP.
DNA encoding human carboxypeptidase.
US2003017574-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-FEB-2003.
(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ) ABX13671 standard; DNA; 55827 BP. 
3 Human protease gene. 
2.8%; Scor
                                                                                                                                                   AAK87551 standard; DNA; 50442 BP.
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Local Similarity 91.2%;
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(APPL-) APPLERA CORP.
  US2002061562-A1.
23-MAY-2002.
(FUKU/) FUKUDA M N.
(AKAM/) AKAMA T O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
RESULT 579
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RESULT 580
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                                                                                                         Local Similarity
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                                                                                        Query Match
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Best Loca RESULT 583

RESULT 582

Length 68355;

Length 86000;

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DE Human cancer associated gene genomic sequence SEQ ID NO:80.

BE Human cancer associated gene genomic sequence SEQ ID NO:80.

BY WOZOGA 6704321-A2.

PD 02-SEP-200474321-A2.

PA (SAGR.) SAGRES DISCOVERY INC.

Query Match

BEST Local Similarity 91.2%; Pred. No. 8.3e-07;

RESULT 595

ID ADG70184 standard; DNA; 379552 BP.

DE DNA of EAC DA236m15-00303.

PD 03-JAN-2003.

PA (Terr.)
                                                                                                                                                                                                                                                                                                                                       PD 03-JAN-2003.
PA (ISIS-) ISIS INNOVATIONS LTD.
Query Match
Best Local Similarity 91.2%; Pred. No. 8.4e-07;
RESULT 596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.8%; Score 68.8; DB 13; Length 113604; 91.2%; Pred. No. 8.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 68.8; DB 10; Length 110000;
Pred. No. 8.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 110000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    'Match 2.8%; Score 68.8; DB 13; Length 113604;
Local Similarity 91.2%; Pred. No. 8.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
2.8%; Score 68.8; DB 8; Length 118951;
Best Local Similarity 91.2%; Pred. No. 8.8e-07;
ID ADQ59167 et-ale.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 2.8%; Score 68.8; DB 9; Length 113604; Best Local Similarity 91.2%; Pred. No. 8.6e-07; RESULT 599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 113604;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADS19853 standard; DNA; 113604 BP.
Human chromosome 21 nucleotide sequence SEQ ID NO:1.
WO2004018701-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADS19854 standard; DNA; 113604 BP.
Human chromosome 21 nucleotide seguence SEQ ID NO:2.
WO2004018701-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.8%; Score 68.8; DB 13; 91.2%; Pred. No. 8.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 2.8%; Score 68.8; DB 9; Best Local Similarity 91.2%; Pred. No. 8.6e-07; RESULT 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABTIT385 standard; DNA; 118951 BP.
Human IG gene related nucleic acid SEQ ID No 11.
WQ200299040-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human LQTS genomic DNA; 113604 BP. US2003077633-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACF42745 standard; DNA; 354391 BP.
Human ALMS1 genomic DNA sequence.
WO2003034072-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADA42607 standard; DNA; 113604 BP
Human LQTS genomic DNA #2.
US2003077633-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 O 04-MAR-2004.
A (PERL-) PERLECEN SCI INC.
A (COXD/) COX D R.
A (ARNO/) ARNOLD D A. 2.84
Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-MAK-20U4.
(PERL-) PERLEGEN SCI INC.
(COXD/) COX D R.
(ARNO/) ARNOLD D A.
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(PERL-) PERLEGEN SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-APR-2003. (PERL-) PERLEGEN SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-APR-2003.
(UYSO-) UNIV SOUTHAMPTON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity RESULT 601
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Best Local Similarity
RESULT 598
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RESULT 602
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ABD33104 standard; DNA; 130877 BP.

Human cancer-associated (CA) gene HD07-009.

N W02004058146-A2.

15-JUL-2004.

A (SAGRE) DISCOVERY INC.

Query Match

2.8%; Score 68.8; DB 13; Length 130877;

Best Local Similarity 91.2%; Pred. No. 9.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ID ADJ37148 standard; cDNA; 139389 BP.

ID ADJ37148 standard; cDNA; 139389 BP.

DE Human malignant pleural mesothelioma (MPM) cDNA #27.

PN US2003219760-Al.

PD 27-NOV-2003.

PA (BGHM) BRIGHAM & WOMENS HOSPITAL INC.

Querry Match 2.8%; Score 68.8; DB 12; Length 139389;

Best Local Similarity 91.2%; Pred. No. 9.6e-07;
                                                                                                                                                                   2.8%; Score 68.8; DB 12; Length 120670; 91.2%; Pred. No. 8.9e-07;
                                                                                                                                                                                                                                          ADQ80254 standard; cDNA; 127145 BP.
HermansKry-Pudlak syndrome associated cDNA.
WO2004063709-A2.
29-JUL-2004.
(BRIM) BRISTCL-MYERS SQUIBB CO.
ery Match
st Local Similarity 73.9%; Pred. No. 9.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.8%; Score 68.8; DB 12; Length 170245; 91.2%; Pred. No. 1.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABK84795 standard; cDNA; 139389 BP.
Human cDNA differentially expressed in granulocytic cells #1366
WO200228999-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 139389;
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Renal cell carcinoma differentially expressed gene #322.
WO2004048933-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAD02697 standard; DNA; 160552 BP.
Human glycosyl sulfotransferase-4 (GST-4) genomic DNA
WO200106015-A1.
MSI-H carcinoma genomic DNA sequence SEQ ID NO:4.
KR2004008012-A.
28-JAN-2004.
(KIMH) KIM H G.
(KIMM) KIM N G.
(LEEJ/) LEE J S.
(RHEE/) RHEE H S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          vuery Match 2.8%; Score 68.8; DB 9; Best Local Similarity 91.2%; Pred. No. 9.6e-07; RESULT 608
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4.
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(GENE-) GENE LOGIC INC.
2.8%; Score 68.8; DB 6;
Best Local Similarity 91.2%; Pred. No. 9.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PD 25-JAN-2001.
PA (REGC) UNIV CALIFORNIA.
Query Match
Best Local Similarity 91.2%; Pred. No. 1e-06;
RESULT 610
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BGHM ) BRIGHAM & WOMENS HOSPITAL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADB70369 standard; cDNA; 139389 BP
PAC 6802 cDNA SEQ ID NO:61.
WO2003021229-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACF62750 standard; DNA; 186591 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (AMHP) WYETH.
(TWAIN) TWINED N C.
(BURC/) BURCZYNSKI M E
(TREP/) TREPICCHIO W L
(DORN/) DORNER A.
(STOV/) STOVER J A.
(SLON/) SLONI D K.
                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity RESULT 605
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
RESULT 611
                                                                                                                                                                                         Best Local Similarity RESULT 604
                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                        Query Match
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20-FEB-2003.
(RPID-) EPIDAUROS BIOTECHNOLOGIE AG.
ery Match 2.8%; Score 68.8; DB 10; Length 208648;
                                                                2.8%; Score 68.8; DB 10; Length 208648; 91.2%; Pred. No. 1.2e-06;
                                                                                                                                                                                                                (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
ry Match 2.8%; Score 68.8; DB 10; Length 208648; t Local Similarity 91.2%; Pred. No. 1.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                            DB 10; Length 208648;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.8%; Score 68.8; DB 10; Length 208648; 91.2%; Pred. No. 1.2e-06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 227968;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABK83497 standard; cDNa; 227968 BP.
Human cDNA differentially expressed in granulocytic cells #68.
WO200228999-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human soft tissue sarcoma-upregulated DNA - SEQ ID 1357
WO2004048938-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 68.8; DB 11;
Pred. No. 1.2e-06;
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Pred. No. 1.3e-06;
                                                                                                                                                                                                                                                                                                                                                                                                          2.8%; Score 68.8; DB 10
91.2%; Pred. No. 1.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADB92113 standard; DNA; 208648 BP.
Human MDR1 related DNA sequence SEQ ID NO:663.
W0200313535-A2.
(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                                                                                                                                                                                                                                                   Human MDR1 related DNA sequence SEQ ID NO:663. WO20031557-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADB96927 standard; DNA; 208648 BP.
Human MDR1 related DNA sequence SEQ ID NO:668.
WO2003013537-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA; 208648 BP.
DNA sequence SEQ ID NO:668
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Human cancer-associated (CA) gene HD07-067.
WO2004058146-A2.
                                                                                                          Human UGTIAl gene sequence SEQ ID NO:663.
20-PPP. A.C.
                                                                                                                                                                                                                                                                                                                                                                20-FEB-2003.
(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                          20-FEB-2003.
(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
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Human genomic sequence hCG41574.
WO2003073826-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PROT-) PROTEIN DESIGN LABS INC.
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11-APR-2002.
(GENE-) GENE LOGIC INC.
2.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 625
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                                                                                    Best Local Similarity
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Best Local Similarity
RESULT 622
                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
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Human MDR1 related
WO2003013535-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
      WO2003013536-A2.
                                                                                                                                                                                            20-FEB-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .Match 2.8%; Score 68.8; DB 10; Length 186591; Local Similarity 91.2%; Pred. No. 1.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 186591;
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                                                                                      Length 186591;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 208648
Cancer based on CYP3A5 related polynucleotide SEQ ID NO:682. WO2003013534-A2.
20-FEB-2003.
(BPID-) EPIDAUROS BIOTECHNOLOGIE AG.
2.8%; Score 68.8; DB 8; Length 1865 of Local Similarity 91.2%; Pred. No. 1.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ACF62735 standard, DNA, 208648 BP.
Cancer based on CYP3A5 related polynucleotide SEQ ID NO:663
WO2003013534-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACF62740 standard, DNA, 208648 BP.
Cancer based on CYP3A5 related polynucleotide SEQ ID NO:668
WO2003013534-A2.
                                                                                                                                           ADB20869 standard; DNA; 186591 BP.
MRP1 based cancer related nucleic acid SEQ ID NO:682.
WO2003013533-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB20850 standard; DNA; 208648 BP.
MRP1 based cancer related nucleic acid SEQ ID NO:663
WO2003013533-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MRP1 based cancer related nucleic acid SEQ ID NO:668.
WO2003013533-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                              2.8%; Score 68.8; DB 10; 91.2%; Pred. No. 1.1e-06;
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                                                                                                                                                                                                         20-FEB-2003.
(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
ery Match 2.8%; Score 68.8; DB 8;
ery Match 91.2%; Pred. No. 1.le-06;
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(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
uery Match
2.8%; Score 68.8; DB 8;
est Local Similarity 91.2%; Pred. No. 1.2e-06;
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Local Similarity 91.2%; Pred. No. 1.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.8%; Score 68.8; DB 8; 91.2%; Pred. No. 1.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADB96941 standard; DNA; 186591 BP.
Human MDR1 related DNA sequence SEQ ID NO:682.
WO200313537-A2.
(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB92132 standard; DNA; 186591 BP.
Human MDR1 related DNA sequence SEQ ID NO:682.
WO2003013535-A2.
                                                                                                                                                                                                                                                                                                                    ADBB7958 standard; DNA; 186591 BP.
Human UGT1A1 gene sequence SEQ ID NO:682
WO2003013536-A2.
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Human UGT1A1 gene sequence SEQ ID NO:668.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-FEB-2003.
(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
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(EPID-) BPIDAUROS BIOTECHNOLOGIE AG
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                                                                                      Query Match
Best Local Similarity
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Best Local Similarity
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Best Local Si
RESULT 613
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Best Local Si
RESULT 617
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Best I RESULT (

Length 213040;

RESULT 618

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Best Local Similarity 90.1%; Pred. No. 5.6e-08; RESULT 638
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                                                                                                                                                                                                                                                                                                                                                                                        Lu 02-JAN-2003.

FU 02-JAN-2003.

PA (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.

PA (ADSC-) CENT ADVANCED                                                                                   vuery watch 2.8%; Score 68.8; DB 13; Length 243390; Best Local Similarity 91.2%; Pred. No. 1.3e-06; RESULT 630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 138627;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 461;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
Query Match 2.8%; Score 68.2; DB 5; Length 415;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 405;
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Novel human diagnostic and therapeutic gene #1837.
WOZOO166753-A2.
13-SEP-ZOO1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.8%; Score 68.6; DB 12; 77.6%; Pred. No. 1.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.8%; Score 68.4; DB 9; 92.3%; Pred. No. 5.2e-08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.8%; Score 68.2; DB 90.1%; Pred. No. 5e-08;
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Human prostate expression marker cDNA 19268.
WO200160860-A2.
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(MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAL61326 standard; DNA; 91000 BP.
Human farnesoid X receptor (FXR) DNA #2.
WO2003044167-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 77.6%; Score RESULT 633

ID ACH27522 standard; CDNA; 461 BP. DE Human adult ovary CDNA #5902. PP UGS203073623-A1. PD 17-APR-2003. PA (IABA/) DRMANG R T. PA (IABA/) IABAT I. PA (IABA/) LABAT I. PA (IABA/) LABAT I. PA (IABA/) LOCAS CONS. PA (IABA/) LOCAS CONS. PA (IABA/) LOCAS CONS. PA (IOCK/) DICKSON M C. PA (JONE/) JONES L W.
                                                                                                                                                                                                                      AAS43104 standard; DNA; 325791 BP. Human Oestrogen receptor beta gene WO200162793-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-JUL-2004.
(SAGR-) SAGRES DISCOVERY INC.
Query Match
                                                                              (SAGR-) SAGRES DISCOVERY INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PA (DRWA) DRWANAC R T.
PA (LABA) LABAT I.
PA (STAC/) STACHE-CRAIN B.
PA (DICK/) DICKSON M C.
PA (JONE/) JONES I. W. 2.8
Query Match 2.8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-MAY-2003.
(ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
RESULT 634
ID AAL61326 standard, D)
DE Human farnesoid X ree
PN WO2003044167-A2.
PD 30-MAY-2003.
PA (ISIS-) ISIS PHARM ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CHIR ) CHIRON CORP. (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                       30-AUG-2001.
(PEKE ) PE CORP NY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
RESULT 635
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Best Local Si
RESULT 636
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AAK85658 standard; DNA; 963 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:40470.
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Pred. No. 1.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-JAN-2003.
(NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.
(ADSC-) CENT ADVANCED SCI & TECHNOLOGY INCUBATIO.
ery Match 2.8%; Score 68.2; DB 10; Length 34118;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 34875
                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.8%; Score 68.2; DB 11; Length 1938; 90.1%; Pred. No. 1.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 5370;
                                                                                                          Length 543;
                                                                                                                                                                                                                                                                                Length 963;
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Melanin-concentrating hormone receptor 1 locus clone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.8%; Score 68.2; DB 10; 90.1%; Pred. No. 6.5e-07;
                                                                                               Query Match 2.84; Score 68.2; DB 4; Best Local Similarity 71.44; Pred. No. 6.5e-08; RESULT 639
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(MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
ery Match
2.8%; Score 68.2; DB 5;
ery Match
7-ncal Similarity 90.1%; Pred. No. 2.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADD14677 standard; cDNA; 34875 BP.
Human src biomarker polynucleotide SEQ ID NO:71.
WQ2003062395-A2.
                                                                                                                                                                                                                                                                                2.8%; Score 68.2; DB 4; 90.1%; Pred. No. 8.9e-08;
AAH12635 standard; cDNA, 543 BP.
Human cDNA clone (3'-primer) SEQ ID NO:9470.
BP1074617-A2.
(HELI-) HELLIX RES INST.
                                                                                                                                                                                                                                                                                                                                            ADM01714 standard; cDNA; 1938 BP.
Human cDNA of the invention SEQ ID NO:399.
EP1347046-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADR52965 standard; DNA; 34875 BP.
Drug therapy altered expressed gene #316.
WO2004072265-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADL45944 standard; DNA; 5370 BP.
Human ovarian cancer DNA marker #19834.
WO200170979-A2.
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Human genomic sequence hCG28567.
WO2003073826-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADC86456 standard; DNA; 34118 BP.
Human GPCR gene SEQ ID NO:909.
EP1270724-A2.
                                                                                                                                                                                                                                                                                                                                                                                                   24-SEP-2003.
(REAS-) RES ASSOC BIOTECHNOLOGY.
Query Match 2.8%; Sco
Best Local Similarity 90.1%; Pre
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(BRIM ) BRISTOL-MYERS SQUIBB CO.
                                                                                                                                                                                                                                      09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
Query Match 2.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (SAGR-) SAGRES DISCOVERY.

ry Match
Local Similarity 90.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (AMHP ) WYETH.
(BURC/) BURCZYNSKI M.
(TWIN/) TWINE N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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Best Local Similarity
                                                                                                                                                                                                                                                                                               Best Local Similarity RESULT 640
                                                                                                                                                                                                                  WO200157182-A2.
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Score 67.8; DB 6; Length 25000;
Pred. No. 7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                     Length 103747;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 318;
                                     Length 5785;
                                                                                                                                                                                                                   Length 6799;
                                                                                                                                                                                                                                                                            ABQ88139 standard; cDNA; 103747 BP.
Human osteoblast differentiation related cDNA SEQ ID NO
WO200250301-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAS27823 standard; DNA; 16489 BP.
DNA encoding novel signal transduction pathway protein,
WO200154733-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABL64391 standard; cDNA; 318 BP.
Human ovarian cancer related cDNA clone SEQ ID NO:7369
WO200192581-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABL68348 standard; DNA; 25000 BP.
Kidney cancer related gene sequence SEQ ID NO:6685.
WO200194629-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 67.8; DB 10;
Pred. No. 5.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           / Match 2.8%; Score 67.8; DB 12; Local Similarity . 91.1%; Pred. No. 2.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.8%; Score 67.8; DB 6;
82.1%; Pred. No. 6.2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.8%; Score 67.8; DB 4;
82.1%; Pred. No. 5.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.8%; Score 67.8; DB 4; 91.1%; Pred. No. 2.6e-07;
                                   Score 68; DB 12;
Pred. No. 2.7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                   Score 68; DB 6;
Pred. No. 1.4e-06;
                                                                                                                                                                                                                 2.8%; Score 68; DB 6; 75.0%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAI63976 standard; DNA; 4096 BP.
Human polynucleotide SEQ ID NO 348.
WO200155308-A2.
                                                                                                               Human gene sequence #17.
W0200252005-A1.
W0200252005-A1.
(KAZU-2002.
(KAZU-3) KAZUSA DNA RES INST FOUND.
(CELE-) CELESTAR LEXICO-SCI LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel human protein DNA #235. 16489 BP. 18200216218.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADM24691 standard; cDNA; 4096 BP.
Human PRO polynucleotide #174.
US2004014039-A1.
                                                                                                BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                              ABN83970 standard; DNA; 6799
                                                                                                                                                                                                                                                                                                                                                                                                   2.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HUMA-) HUMAN GENOME SCI INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.8%;
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                                                      80.08;
                                                                                                                                                                                                                                                                                                                                    27-JUN-2002.
(GENE-) GENE LOGIC INC.
(PROC ) PROCTER & GAMBLE CO.
                                     2.88;
                  GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AVAL-) AVALON PHARM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-DEC-2001.
(CORI-) CORIXA CORP.
                                                        Best Local Similarity
                                                                                                                                                                                                                                      Local Similarity
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(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
RESULT 664
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Local Similarity
08-APR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-DEC-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                     Query Match
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                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                     Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 663
                                                                                                                                                                                                                                                                                                                                                                                                       Query
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADJ09023 standard; DNA; 3560 BP.
Human cardiovascular system associated polypeptide-related DNA SeqID2411.
US2004005575-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADJ09022 standard; DNA; 3561 BP.
Human cardiovascular system associated polypeptide-related DNA SeqID2410.
US2004005575-A1.
                                                        2.8%; Score 68.2; DB 12; Length 122557; 90.1%; Pred. No. 1.3e-06;
                                                                                                                                                                                                                     Length 247682;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAS36910 standard; DNA; 3561 BP.
Human cardiovascular system antigen genomic DNA SEQ ID No 2410.
WC200155321-A2.
02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
2.8%; Score 68; DB 4; Length 3561;
st Local Similarity 80.0%; Pred. No. 2.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human cardiovascular system antigen genomic DNA SEQ ID No 2411 WO200155321-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 3561;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 3560;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 3561;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 3560
                                                                                                                                                                                                                                                                                                                                                                                    Length 1812;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 3560;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADE47605 standard; DNA; 3560 BP.
Human cardiovascular system related genomic DNA #1171.
US2003059908-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADE47604 standard; DNA; 3561 BP.
Human cardiovascular system related genomic DNA #1170.
US2003059908-A1.
                                                                                                                                                                                                       vuery Match 2.8%; Score 68.2; DB 12;
Best Local Similarity 90.1%; Pred. No. 1.9e-06;
RESULT 648
                                                                                                                                                                                                                                                                                                                                07-FEE-2001.
(HELI-) HELIX RES INST.
(ery Match 2.8%; Score 68; DB 4; I
2.8%; Score 68; DB 4; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             O2-AUG-2001.
(HUWAL) HUWAN GENOME SCI INC.
ery Match
2.8%; Score 68; DB 4; I
2.8%; Score 68; DB 4; I
2.8%; Pred. No. 2.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-MAR-2003.

(HUMA-) HUMAN GENOME SCI INC.

Query Match

2.8%; Score 68; DB 10;

Best Local Similarity 80.0%; Pred. No. 2.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-2x1-2004.

08-3x1-2004.

(HUMA-) HUMAN GENOME SCI INC.

2.8%; Score 68; DB 13;

ery Match

c', milarity 80.0%; Pred. No. 2.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-MAR-2003.
(HUMA-) HUMAN GENOME SCI INC.
Ery Match 2.8%; Score 68; DB 10;
ery Match 6.00%; Pred. No. 2.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-JAN-2004.
(HUMA-) HUMAN GENOME SCI INC.
r: Watch
tt Local Similarity 80.0%; Pred. No. 2.1e-07;
                                                                                              Human gene associated with low HDL-C AT3. US204043389-A1.
                                                                                                                                                                                                                                                                   AAH18046 standard; CDNA; 1812 BP.
Human CDNA sequence SEQ ID NO:17874.
EP1074617-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADNO6010 standard; cDNA; 5785 BP. Antipsoriatic cDNA sequence #1240.WO2004028479-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA; 3560 BP.
                  18-DEC-2003.
(UYPH-) UNIV PHILIPPS MARBURG.
                                                                                                                                                                             04-MAR-2004.
(VITI-) VITIVITY INC.
                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 649
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Matcn
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 650
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAS36911 standard;
WO2003104489-A2.
                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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Seq ID 1483

Length 16489;

RESULT 653

Length 4096;

Length 4096

46.

Length 16489;

Length 110000

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2.8%; Score 67.6; DB 10; Length 125515; 72.9%; Pred. No. 1.9e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAK89420 standard, DNA, 185 BP.
Human digestive system antigen genomic sequence SEQ ID NO: 2996.
WO200155314-A2.
                                                                                                                                                                                                                                                                                                                           Human digestive system antigen genomic sequence SEQ ID NO: 2995.
WO200155314-A2.
                Score 67.6; DB 6; Length 36741; Pred. No. 9.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-4PR-2003.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.4; DB 11; Length 185;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match (1987) Score 67.4; DB 4; Length 185;
ery Match (1981) 87.1%; Pred. No. 5.9e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABN90280 standard; DNA; 185 BP.
Human liver antigen HLDR194 genomic sequence, SEQ ID NO:401.
US2002042096-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABN90279 standard; DNA; 185 BP.
Human liver antigen HLDR194 genomic sequence, SEQ ID NO:400
US2002042096-Al.
                                                                                                                                       ADL13941 standard; DNA; 125515 BP.
Ostcoarthritis-associated polymorphic nucleotide #473.
WO2001054166-A2.
(13-JUL-2003.
(INCY-) INCYTE GENOMICS INC.
                                                                              2.8%; Score 67.6; DB 10; 73.5%; Pred. No. 1.8e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HUMA-) HUMAN GENOME SCI INC.
ry Match
L Local Similarity 87.1%; Pred. No. 5.9e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 67.4; DB 6;
Pred. No. 5.9e-08;
                                                                                                                                                                                                                                                                                                                                                             02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
(HTMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.4; DB 4;
lery Match 2.7%; Pred. No. 5.9e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.7%; Score 67.4; DB 6;
87.1%; Pred. No. 5.9e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.7%; Score 67.4; DB 5;
87.1%; Pred. No. 5.9e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADJ15193 standard; DNA; 185 BP.
Human liver-related genomic DNA - SEQ ID 401.
US2003077602-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAS31925 standard; DNA; 185 BP.
Human liver associated genomic DNA #99.
WO200155355-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAS31924 standard; DNA; 185 BP.
Human liver associated genomic DNA #98.
WO200155355-A1.
                                                                                                                                                                                                                                                                                                          AAK89419 standard; DNA; 185 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.7%;
                     2.8%;
(TEXA ) UNIV TEXAS SYSTEM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
                                                                                                                                                                                                                                                            Best Local Similarity RESULT 676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity RESULT 679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-APR-2002.
(ROSE/) ROSEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ROSE/) ROSEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity RESULT 682
                Query Match
Best Local Similarity
RESULT 674
                                                                            Query Match
Best Local Similarity
RESULT 675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 680
                                                                                                                                                                                                                                                                                                                                                                                                                Query
                Human immune/haematopoietic antigen genomic sequence SEQ ID NO:31915.
W0200157182-A2.
09-AUG-2001.
09-AUG-2001.
Local Similarity 82.1%; Pred. No. 7.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 110000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 127722;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              vuery match 2.8%; Score 67.6; DB 6; Length 36741;
Best Local Similarity 89.0%; Pred. No. 9.8e-07;
RESULT 672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 36741;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 36741;
                                                                                                                                                                                                                                                                    Length 27571;
                                                                                                                                                                                                                                                                                                                                                                                                                               Length 27572;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 826;
                                                                                                                                                                         Human nervous system related polynucleotide SEQ ID NO 11553
                                                                                                                                                                                                                                                                                                                           ABA19221 standard; DNA; 27572 BP.
Human nervous system related polynucleotide SEQ ID NO 11552
WO200159063-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AA196289 standard; cDNA; 826 BP.
Human neuroblastoma expressed polynucleotide SEQ ID NO 2364
WO200166719-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABQ88153 standard; cDNA; 36741 BP.
Human osteoblast differentiation related cDNA SEQ ID NO 60.
WO20025031-A2.
27-JUN-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADQ97301 standard; DNA; 127722 BP.
Human cancer associated sequence HD08-028, SEQ ID 278.
WO2004060304-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.8%; Score 67.8; DB 12;
91.1%; Pred. No. 1.6e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22-JUL-2004.
(SAGR-) SAGRES DISCOVERY INC.
LY MARCH
LOCAL Similarity 82.1%; Pred. No. 1.7e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WOZOULUNDO

13-SEP-2001.

(CHE) CHIBA PREFECTURE.

(HISM ) HISAMITSU PHARM CO LTD.

2.8%; Score 67.6; DB 4;
                                                                                                                                                                                                                                                                                                                                                                                   16-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match
2.8*; Score 67.8; DB 5;
ery Match
91.1*; Pred. No. 7.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 67.6; DB 4;
Pred. No. 9.8e-07;
                                                                                                                                                                                                                    16-AUG-2001.

(HUMA-) HUMAN GENOME SCI INC.

2.8%; Score 67.8; DB 5;

ery Match 2.8%; Pred. No. 7.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (UTNY) UNIV NEW YORK STATE RES FOUND.
ry Match
t Local Similarity 89.0%; Pred. No. 9.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABS52806 standard; DNA; 36741 BP.
DNA encoding human adenosine deaminase (ADA).
US2002088017-A1.
04-JUL-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAS03388 standard; DNA; 36741 BP.
Human adenosine deaminase (ADA) genomic DNA
US6207876-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABS66890 standard; DNA; 36741 BP.
Human adenosine deaminase (ADA) gene.
US2002102731-Al.
                                                                                                                                                            ABA19222 standard; DNA; 27571 BP.
  AAK77103 standard; DNA; 26372 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PD 27-MAR-2001.
PA (TEXA) UNIV TEXAS SYSTEM.
Query Match
Best Local Similarity 89.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A (GENE-) GENE LOGIC INC.
A (PROC ) PROCTER & GAMBLE CO.
Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Match 2.8%;
Local Similarity 89.0%;
                                                                                                 Query Match
Best Local Similarity
RESULT 665
ID ABA1922 standard; DN
DE Human nervous system
PN W020159063-A2.
PD 16-AUG-2001.
PA (HUMA-) HUMAN GENOME
                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
REGULT 666
ID ABA1921 standard, D)
DE Human nervous system
PN WO20015965-A2.
PD 16-AUG-2001.
PA (HUMA-) HUMAN GENOME
                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
RESULT 667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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Best Loc.
RESULT 670
ID AAS03:
DE Hume.
PD 7
PD 7
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Length 185;

Length

Length 185;

Length 185

Length 185;

24-APR-2003

Query Match

Query Match

RESULT 685

Query Match

Query Match

Best Loca RESULT 688

Query Match

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AAF21005 standard, DNA; 4704 BP.
Human low adenosine antisense oligonucleotide related seguence #2572.
WO200062736-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABK84729 standard; cDNA; 4704 BP.
Human cDNA differentially expressed in granulocytic cells #1300.
WO200228999-A2.
                                                                                                                                                                                                                                              AAA34883 standard; DNA; 4704 BP.
Human adenosine receptor related polynucleotide SEQ ID NO:2572.
WO200009525-A2.
24-FEB-2000.
                     2.7%; Score 67.4; DB 12; Length 4654; 82.8%; Pred. No. 3.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.7%; Score 67.4; DB 10; Length 4704; 82.8%; Pred. No. 3.6e-07;
                                                                                                                                                                                    Length 4698;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.7%; Score 67.4; DB 11; Length 4704; 82.8%; Pred. No. 3.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 4704;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 4704;
                                                                                                                                                                                                                                                                                                                                                    Length 4704;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 4704;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 4704;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADQ22218 standard; DNA; 4800 BP.
Human soft tissue sarcoma-upregulated DNA - SEQ ID 5038.
WQ2004048938-A2.
10-JUN-2004.
                                                                                ADN01990 standard; DNA; 4698 BP.
Human inflammatory bowel disease marker gene - CD11c.
JP2004065120-A.
                                                                                                                                        04-MAR-2004.
(SUMU) SUMITOMO SEIYAKU KK.
ery Match 2.7%; Score 67.4; DB 12;
st Local Similarity 82.8%; Pred. No. 3.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADR44042 standard; DNA; 4704 BP.
Human kidney tumour associated gene clone-53 SEQ
WO2004074506-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.7%; Score 67.4; DB 11;
82.8%; Pred. No. 3.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 67.4; DB 13;
Pred. No. 3.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABD20548 standard; DNA; 4704 BP.
Human pulmonary and inflammatory target DNA #159.
WO200285309-A2.
                                                                                                                                                                                                                                                                                                                                                    Score 67.4; DB 3;
Pred. No. 3.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 67.4; DB 3;
Pred. No. 3.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 67.4; DB 6;
Pred. No. 3.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADI32032 standard; cDNA; 4704 BP.
Human cDNA #1358.
    (PROT-) PROTEIN DESIGN LABS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA; 4704 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-OCT-2002.
(EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                      2.7%;
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(EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human nucleic acid sequence. WO200285308-A2.
                                                                                                                                                                                                                                                                                                                                 (UYEC-) UNIV EAST CAROLINA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYEC-) UNIV EAST CAROLINA.
(NYCE/) NYCE J W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-APR-2002.
(GENE-) GENE LOGIC INC.
                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MERG-) MERGEN LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABZ96699 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US6607879-B1.
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                                                                                                                                                                                       Query Match
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RESULT
                                                                                                                                                                                                                                                                                                                                                                                                                AAK68070 standard; DNA; 2480 BP.

Human immune/haematopoietic antigen genomic sequence SEQ ID NO:22882.
W0200157182-A2.
90-5001-7182-A2.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.4; DB 4; Length 2480;
ery Match
Et Local Similarity 87.1%; Pred. No. 2.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAF21006 standard; DNA; 4654 BP.
Human low adenosine antisense oligonucleotide related sequence #2573.
WO200062736-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA34884 standard; DNA; 4654 BP.
Human adenosine receptor related polynucleotide SEQ ID NO:2573.
WO200009525-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.7%; Score 67.4; DB 11; Length 2884; 87.1%; Pred. No. 2.7e-07;
                                                                                                                                              Length 185;
                                                                                                                                                                                                                                                                                                                                                      Query Match 2.7%; Score 67.4; DB 10; Length 583; Best Local Similarity 87.1%; Pred. No. 1.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 4654;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31-OCT-2002.
(BPIG-) EPIGENESIS PHARM INC.
ery Match 2.7%; Score 67.4; DB 11;
ery Match 82.8%; Pred. No. 3.5e-07;
                                                                                                                       (HUMA-) HUMAN GENOME SCI INC.

ry Match

t Local Similarity 87.1%; Pred. No. 5.9e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.7%; Score 67.4; DB 3;
82.8%; Pred. No. 3.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABD20549 standard, DNA, 4654 BP.
Human pulmonary and inflammatory target DNA #160.
WO200285309-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 67.4; DB 3;
Pred. No. 3.5e-07;
    87.1%; Pred. No. 5.9e-08;
                                                                                                                                                                                                 ACD96444 standard; cDNA; 583 BP.
Human colon cancer cell expressed cDNA #4856.
US2002155438-Al.
                                          ADJ15192 standard; DNA; 185 BP.
Human liver-related genomic DNA - SEQ ID 400.
US2003077602-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADM03547 standard; cDNA; 2884 BP.
Human cDNA of the invention SEQ ID NO:2232.
EP1347046-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA; 4654 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADQ17509 standard; DNA; 4654 BP.
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24-FEB-2000.
(UYEC-) UNIV EAST CAROLINA.
ery Match 2.7%;
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(EPIG-) EPIGENESIS PHARM INC.
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                                                                                                                                                                                                                                                               24-0CT-2002.
(SIMP/) SIMPSON A J G.
(NETO/) NETO E D.
(BREN/) BRENTANI R R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity RESULT 690
Best Local Similarity
RESULT 683
                                                                                                                                                             Best Local Similarity
RESULT 684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 686
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RESULT 687
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10-JUN-2004

Query Match

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(HUMA-) HUMAN GENOME SCI INC.
    02-AUG-2001
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                                                                                       RESULT 710
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                                                                                                                                                                                                                                                        AAF21008 standard; DNA; 11649 BP.
Human low adenosine antisense oligonucleotide related sequence #2575.
WO200062736-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADP49338 standard; DNA; 80000 BP.
Human B-cell chronic lymphocytic leukaemia associated genomic DNA #2.
WO2004046376-A2.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:26887.
WO200157182-A2.
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DNA encoding novel signal transduction pathway protein, Seq ID 1442.
WO200154733-A1.
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PA (UYMC-) UNIV MCMASTER.

Query Match 2.7%; Score 67.4; DB 12; Length 25373;
Best Local Similarity 76.1%; Pred. No. 9.1e-07;
RESULT 706
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 10; Length 11649;
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                                                                                     AAA34886 standard; DNA; 11648 BP.
Human adenosine receptor related polynucleotide SEQ ID NO:2575.
WO200009525-A2.
      QUELY MACCH 2.7%; Score 67.4; DB 12; Length 4800; Best Local Similarity 82.8%; Pred. No. 3.6e-07; RESULT 701
                                                                                                                      PD 24-FEB-2000.
PA (UYEC-) UNIV EAST CAROLINA.
Query Match 2.7%; Score 67.4; DB 3; Length 11648;
Best Local Similarity 82.8%; Pred. No. 5.9e-07;
                                                                                                                                                                                                                                                                                                                                                                                          2.7%; Score 67.4; DB 3; Length 11649; 82.8%; Pred. No. 5.9e-07;
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Human PAZ/PIWI domain-containing protein polynucleotide #3
US2003232442-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOZOCZ-202.
31-OCT-2002.
SI-OCT-2002.
(EPIG-) EPIGENESIS PHARM INC.
2.7%; Score 67.4; DB 11;
ery Match
2.7%; Score 67.4; DB 11;
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(ISIS-) ISIS PHARM INC.
2.74; Score 67.4; DB 12;
FL Local Similarity 92.24; Pred. No. 2.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABD20551 standard; DNA; 11649 BP.
Human pulmonary and inflammatory target DNA #162
WO200285309-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
12. Match 2.7%; Score 67.2; DB 4;
t Local Similarity 90.0%; Pred. No. 4.8e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABZ96702 standard; DNA; 11649 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADK16022 standard; DNA; 25373 BP.
Human mammaglobin gene promoter.
WO2004016652-A2.
(PROT-) PROTEIN DESIGN LABS INC.
                                                                                                                                                                                                                                                                                                                       26-OCT-2000.
(UYEC-) UNIV BAST CAROLINA.
(NYCE/) NYCE J W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HSRI-) HS RIGSHOSPITALET.
(UYAR-) UNIV ARHUS.
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                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity RESULT 703
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Best Local Similarity
RESULT 705
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Best Local Si
RESULT 708
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AAK67157 standard; DNA; 128 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:21969.
WO200157182-A2.
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Genomic sequence #4 encoding for novel human respiratory antigen
WO200155448-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAK89597 standard; DNA; 144 BP.
Human digestive system antigen genomic sequence SEQ ID NO: 3173.
WO200155314-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAK89594 standard; DNA; 157 BP.
Human digestive system antigen genomic sequence SEQ ID NO: 3170.
WO200155314-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAK89595 standard; DNA; 154 BP.
Human digestive system antigen genomic sequence SEQ ID NO: 3171
WO200155314-A2.
                                                                                                                                                                                                                        Score 67.2; DB 10; Length 110;
Pred. No. 5e-08;
2.7%; Score 67.2; DB 4; Length 110; 90.0%; Pred. No. 5e-08;
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Human respiratory system associated genomic DNA seg id 598.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAS34757 standard; DNA; 142 BP.
Human DNA for a novel foetal antigen, SEQ ID No 2181.
WO200155312-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAS34756 standard; DNA; 142 BP.
Human DNA for a novel foetal antigen, SEQ ID No 2180
WO200155312-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
(HTWA-) HUMAN GENOME SCI INC.
(ery Match 2.7%; Score 67.2; DB 5;
(ery Match 5.8e-08;
                                                                                                                                                                                                                                                                                                                                                 09-ACG-2001.
(HUWA-) HUMAN GENOME SCI INC.
(HTWA-) HUMAN GENOME SCI INC.
(ery Match 2.7%; Score 67.2; DB 4;
(ery Match 5.5e-08;
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02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 5;
ery Match
2.7%; Score 67.2; DB 5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.7%; Score 67.2; DB 4; 90.0%; Pred. No. 6.1e-08;
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90.0%; Pred. No. 6.2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.7%; Score 67.2; DB 4; 90.0%; Pred. No. 5.8e-08;
                                                              ADB94585 standard; DNA; 110 BP.
Novel human protein DNA #194.
US2002168711-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
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(HUMA-) HUMAN GENOME SCI INC.
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                                                                                                                                                                                                                        2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                       HESULT 712
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Best Local Similarity
                                                                                                                                  14-NOV-2002.
(ROSE/) ROSEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 714
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
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Query Match
Best Local Similarity
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02-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query
                                                                                                                                                                                                                                                                               Human respiratory system associated polypeptide-related DNA SeqID598.
US2003077704-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAK89421 standard; DNA; 186 BP.
Human digestive system antigen genomic sequence SEQ ID NO: 2997
WO200155314-A2.
US2003215893-A1.
20-NOV-2003.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 10; Length 160; set Local Similarity 90.0%; Pred. No. 6.2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 160
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 186;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-AUG-2001.
(HUMA) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 4; Length 186;
rt Local Similarity 90.0%; Pred. No. 6.7e-08;
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Human liver antigen HLDR194 genomic sequence, SEQ ID NO:402.
US2002042096-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABA16882 standard; DNA; 172 BP.
Human nervous system related polynucleotide SEQ ID NO 9213
WO200159063-A2.
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Human nervous system related polynucleotide SEQ ID NO 8663.
WO200159063-A2.
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PA (HUMA-) HUMAN GENOME SCI INC.
Query Match
2.7%; Score 67.2; DB 11;
RESULT 726
RESULT 726
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(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 67.2; DB 5;
ery Match 90.0%; Pred. No. 6.4e-08;
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Best Local Similarity 90.0%; Pred. No. 6.7e-08;
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ry Match

t Local Similarity 90.0%; Pred. No. 6.7e-08;
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ry Match
t Local Similarity 90.0%; Pred. No. 6.7e-08;
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ID ADJ15194 standard; DNA; 186 BP.
DE Human liver-related genomic DNA - SEQ ID 402.
PN US2003077602-A1.
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Human liver associated genomic DNA #100.
WO200155355-A1.
                                                                                                                                                                                                                                                  DNA; 160 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
RESULT 720
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(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                  ADI97134 standard;
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AAK67132 standard; DNA; 310 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:21944.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAK72194 standard; DNA; 336 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:27006.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABZ74226 standard; DNA; 312 BP.
Secreted protein gene 215 genomic fragment HNGAM58, SEQ ID NO:1373.
WO200277013-A2.
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Human reproductive system related antigen DNA SEQ ID NO: 7791.
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                                                                                                     Length 275;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.7%; Score 67.2; DB 8; Length 312; 90.0%; Pred. No. 9e-08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 310;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABZ67792 standard; DNA; 312 BP.
Human secreted protein encoding genomic DNA SEQ ID NO 1315.
WO200277186-A2.
                                                                                                                                                                                                                                                                                                                                                                    AAZ13311 standard; cDNA; 294 BP.
Human gene expression product cDNA sequence SEQ ID NO:780.
WO9938972-A2.
Human excretory related polynucleotide SEQ ID NO 778. WO200155313-A2.
                                                                                                                                                                       AAI63364 standard; DNA; 275 BP.
Human kidney related polynucleotide SEQ ID NO 679.
WO200155323-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADA98755 standard; DNA; 312 BP.
Human secreted protein-related DNA sequence #348
WO2003004623-A2.
                                                                                                                                                                                                                                                                   (HUMA-) HUMAN GENOME SCI INC.

2.7%; Score 67.2; DB 5;

t Local Similarity 90.0%; Pred. No. 8.4e-08;
                                                                         (HUMA-) HUMAN GENOME SCI INC.

2.7%; Score 67.2; DB 4;

t Local Similarity 90.0%; Pred. No. 8.4e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.7%; Score 67:2; DB 4; 90.0%; Pred. No. 8.9e-08;
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2.7%; Score 67.2; DB 8;
t Local Similarity 90.0%; Pred. No. 9e-08;
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90.0%; Pred. No. 9.3e-08;
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(HUMA-) HUMAN GENOME SCI INC.
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(HUMA-) HUMAN GENOME SCI INC.
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(CHIR ) CHIRON CORP.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                       Hest Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
RESULT 730
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ACA03395 standard; DNA; 424 BP.
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    BABBBBB
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Genomic sequence #18 encoding for novel human respiratory antigen
WO200155448-A1.
                                                                                                                                       PD 02-AUG-2001.

PA (HUMA-) HUMAN GENOME SCI INC.

Query Match
Best Local Similarity 90.0%; Pred. No. 9.3e-08;

RESULT 738

DE Human testicular antigen encoding DNA fragment SEQ ID NO: 2648.

PD 02-AUG-2001.

PA (HUMA-) HUMAN CENOME SCI INC.

2.7%; Score 67.2; DB 4; Length 336;

RESULT 738

DE Human testicular antigen encoding DNA fragment SEQ ID NO: 2648.

PA (HUMA-) HUMAN CENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADD71335 standard; DNA; 360 BP.
Glutamine:fructose-6-phosphate amide transferase 1 gene intron
WO2003023063-A1.
                                                                                                                                   ABL97995 standard; DNA; 336 BP.
Human testicular antigen encoding DNA fragment SEQ ID NO: 2647
WO200155317-A2.
  Human reproductive system related antigen DNA SEQ ID NO: 7790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 355;
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PA (HUMA-) HUMAN GENOME SCI INC.

Query March

Best Local Similarity 90.0%; Pred. No. 1.1e-07;

RESULT 743
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                                                                                                                                                                                                                                                                                                                                                                                               Length 336;
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                                                                               Length 336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAS30632 standard; DNA; 424 BP.
DNA encoding novel lung cancer antigen, Seq ID No WO200155300-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.7%; Score 67.2; DB 10; 90.0%; Pred. No. 9.7e-08;
        PN WO200155320-A2.

PD 02-AUG-2001.

PA (HUMA-) HUMAN GENOME SCI INC.

Query Match

Best Local Similarity 90.0%; Pred. No. 9.3e-08;
                                                                                                                                                                                                                                                                                                                                            W.c.v....
02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 67.2; DB 4;
ery Match 90.0%; Pred. No. 9.3e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                               ACD92142 standard; cDNA; 355 BP.

Human colon cancer cell expressed cDNA #554.

US2002155438-Al.

24-OCT-2002.

(SIMP/) SIMPSON A J G.

(NETO/) NETO E D.

(REEN/) BRENTANI R R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AA192285 standard; cDNA; 424 BP.
Human polynucleotide SEQ ID NO 12345.
WO200164835-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAI86639 standard; cDNA; 394 BP.
Human polynucleotide SEQ ID NO 6699.
WO200164835-A2.
(HYSE-2001.
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(HUMA-) HUMAN GENOME SCI INC.
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A (SANY ) SANKYO CO LTD.
Query Match
Best Local Similarity 90
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Best Local Similarity
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Best Local Similarity
RESULT 739
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Best Local Similarity
RESULT 745
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(HYSE-) HYSEQ INC.
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AD197148 standard; DNA; 424 BP.
Human respiratory system associated polypeptide-related DNA SeqID612.
US2003077704-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 / Match 2.7%; Score 67.2; DB 11; Length 424; Local Similarity 90.0%; Pred. No. 1.1e-07;
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                                                                                                                                               Length 424;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 429;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 451;
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                                                                                                                                                                                                                                                                                                                                                                                                     ADG41374 standard; DNA; 424 BP.

Human respiratory system associated genomic DNA seg id 612.
US2003215893-A1.
20-NOV-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABL87490 standard; cDNA; 451 BP.

Human ovarian cancer related cDNA clone SEQ ID NO:10468.

#0200192581-A2.

06-DEC-2001.

(CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HUMA-) HUMAN GENOME SCI INC.

2.7%; Score 67.2; DB 10;

t Local Similarity 90.0%; Pred. No. 1.1e-07;
                                                                                                                                             2.7%; Score 67.2; DB 8; 90.0%; Pred. No. 1.1e-07;
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Query Match 2.7%; Score 67.2; DB 5;
Best Local Similarity 90.0%; Pred. No. 1.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-AUG-2001.
(MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
2.7%; Score 67.2; DB 5;
cery Match 2.7%; Pred, No. 1.1e-07;
                                                                                                                                                                                                                                                                                                                                  2.7%; Score 67.2; DB 9;
90.0%; Pred. No. 1.1e-07;
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(MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.

Query Match

2.7%; Score 67.2; DB 5;

Best Local Similarity 90.0%; Pred. No. 1.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.7%; Score 67.2; DB 6;
90.0%; Pred. No. 1.1e-07;
DNA encoding human lung cancer antigen HCLSC85.
US2002173454-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABV19126 standard; cDNA; 429 BP.
Human prostate expression marker cDNA 19317.
WO200160860-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABV50841 standard; cDNA; 463 BP.
Human prostate expression marker cDNA 50832.
WO200160860-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABV61721 standard; cDNA; 470 BP.
Human prostate expression marker cDNA 61712.
WO200160860-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC
                                                                                                                                                                                                                ADB96743 standard; DNA; 424 BP.
Novel lung cancer antigen genomic DNA #14
US2003049703-A1.
13-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADL41307 standard; DNA; 491 BP.
Human ovarian cancer DNA marker #15197.
WO200170979-A2.
                                                                                                                                                                                                                                                                                                              (HUMA-) HUMAN GENOME SCI INC.
Query Match 2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24-APR-2003. (HUMA-) HUMAN GENOME SCI INC.
                                            21-NOV-2002.
(ROSE/) ROSEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
RESULT 752
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Best Local Similarity
RESULT 748
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 747
                                                                                                                                          Ouery Match
Best Local Similarity
RESULT 746
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Length 1167;

Length 1297

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AAK70343 standard; DNA; 1389 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:25155.
WO200157182-A2.
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Genomic sequence #359 encoding human colon associated polypeptide.
WO200155302-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genomic sequence #362 encoding human colon associated polypeptide WO200155302-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human digestive system antigen genomic sequence SEQ ID NO: 3962. WO200155314-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAK90385 standard; DNA; 1713 BP.
Human digestive system antigen genomic sequence SEQ ID NO: 3961
WO200155314-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.7%; Score 67.2; DB 6; Length 1440; 90.0%; Pred. No. 2.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                               (HUMA-) HUMAN GENOME SCI INC.

ry Match 2.7%; Score 67.2; DB 4; Length 1389;

t Local Similarity 90:0%; Pred. No. 2.1e-07;
                                                                                                                         AAX24734 standard; DNA; 1297 BP.
Human APP genomic clone containing polyadenylation signals.
WO9909150-A1.
                                                                  2.7%; Score 67.2; DB 4;
90.0%; Pred. No. 1.9e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
(HTWA-) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 4;
ery Match 2.38-07;
                                                                                                                                                                                                                                            Score 67.2; DB 2;
Pred. No. 2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.7%; Score 67.2; DB 6;
90.0%; Pred. No. 2.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.7%; Score 67.2; DB 4; 90.0%; Pred. No. 2.3e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 4;
Local Similarity 90.0%; Pred. No. 2.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SHAN-) SHANGHAI BIOWINDOW GENE DEV INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (BODE-) BODE GENE DEV CO LTD SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human polynucleotide SEQ ID NO 9542. W202016453-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABQ77752 standard; cDNA; 1440 BP.
Serine protease 9.02-encoding cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABS56842 standard; cDNA; 1389 BP.
Human HUMSIAH protein 14.74 cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAK90386 standard; DNA; 1713 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA; 1713 BP
  21-DEC-2000.
(HUMA-) HUMAN GENOME SCI INC.
(ROSE/) ROSEN C A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
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766
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Best Local Similarity
RESULT 764
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                                                                                  Best Local Similarity
RESULT 763
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(FARB ) BAYER CORP.
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Best Local S
                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAK65574 standard; DNA; 845 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:20386.
WO200157182-A2.
                                                                                                                                                                                                                                          Human cDNA sequence useful for the treatment of cancer (SeqID 1071) WO2003050236-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAL37008 standard; DNA; 1130 BP.
Human musculoskeletal system related polynucleotide SEQ ID NO 3373.
WO200155367-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADJ30746 standard; DNA; 1130 BP.
Human musculoskeletal system-associated genomic DNA - SEQ ID 3373.
US2004009488-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABX59996 standard; cDNA; 1130 BP.
cDNA encoding novel human musculoskeletal system antigen #2340.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABA20629 standard; DNA; 1130 BP.
Human nervous system related polynucleotide SEQ ID NO 12960
WO200159063-A2.
RESULT 754

ID AAZ15513 standard; CDNA; 745 BP.

BE Human gene expression product CDNA sequence SEQ ID NO:2982.

BN W09938972-A2.

PD 05-AUG-1999.

PA (CHIR ) CHIRON CORP.

PA (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                        ABQ89819 standard; cDNA; 834 BP.
Human prostate expressed polynucleotide SBQ ID NO 1075
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Match 2.7%; Score 67.2; DB 12;
Local Similarity 90.0%; Pred. No. 1.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 67.2; DB 4;
ery Match 90.0%; Pred. No. 1.8e-07;
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(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 4;
tt Local Similarity 90.0%; Pred. No. 1.6e-07;
                                                                                                                                                       2.7%; Score 67.2; DB 2; 90.0%; Pred. No. 1.5e-07;
                                                                                                                                                                                                                                                                                                                                                          2.7%; Score 67.2; DB 9; 90.0%; Pred. No. 1.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.7%; Score 67.2; DB 6; 90.0%; Pred. No. 1.5e-07;
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90.0%; Pred. No. 1.8e-07;
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Human secreted protein gene 24 SEQ ID NO:34.
WO200077026-A1.
                                                                                                                                                                                                                         ADB82858 standard; cDNA; 763 BP
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(HUMA-) HUMAN GENOME SCI INC.
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(HUMA-) HUMAN GENOME SCI INC.
                                                                                    05-AUG-1999.
(CHIR ) CHIRON CORP.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                            (CHIR ) CHIRON CORP. (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CHIR ) CHIRON CORP. (HYSE-) HYSEQ INC.
                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ROSE/) ROSEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
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Length 1674;

Length 1389;

RESULT 757

Length 1713;

Length 1713

Length 1713

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Query Match
Best Local Similarity
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Full length human cDNA useful for treating neurological disease Seq 1295.
EP1447413-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cDNA encoding a human transporter & ion channel (TRICH) protein SeqID125. W02003093444-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human secreted protein-encoding gene 10 cDNA clone HCEGK81, SEQ ID NO:20. WO200077022-A1.
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Human cDNA sequence useful for the treatment of cancer (SeqID 1459).
WO2003050236-A2.
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(HUWA-) HUMAN GENOME SCI INC.
ery Match imilarity 2.7%; Score 67.2; DB 6; Length 2246;
                                          Length 1713;
                                                                                                                                                                                                                                                           2.7%; Score 67.2; DB 9; Length 1713; 90.0%; Pred. No. 2.3e-07;
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                                                                                                                              837.
                                                                                                                                                                                                                                                                                                                     ADB32903 standard; DNA; 1713 BP.
Human novel colon related polypeptide DNA SEQ ID NO 840.
US2003050231-Al.
                                                                                                    ADB12900 standard; DNA; 1713 BP.
Human novel colon related polypeptide DNA SEQ ID NO
US2003050231-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAS19209 standard; cDNA; 1873 BP.
DNA encoding human transformer 2-beta protein 29.15.
W0200192517-A1.
06-DEC-2001.
(SHAN-) SHANGHAI BIOWINDOW GENE DEV INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PD 18-AUG-2004.

PA (REAS-) RES ASSOC BIOTECHNOLOGY.

Query Match

2.7%; Score 67.2; DB 13;

Best Local Similarity 90.0%; Pred. No. 2.6e-07;

RESULT 779
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      PA (HUMA-) HUMAN GENOME SCI INC.
Query Match
2.7%; Score 67.2; DB 5;
Best Local Similarity 90.0%; Pred. No. 2.3e-07;
RESULT 772
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-DEC-2000.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 67.2; DB 4;
ery Match 2.7%; Pred. No. 2.4e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CDNA; 1786 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADH22627 standard; cDNA; 1774 BP
                                                                                                                                                                  13-MAR-2003.
(ROSE/) ROSEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (INCY-) INCYTE CORP. Query Match
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                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
RESULT 773
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
RESULT 775
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 774
                                                                                                                                                                                                                                                                                                                                                                                                       (ROSE/) ROSEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAD08354 standard;
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02-AUG-2001
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Best Local Si
RESULT 777
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Tumour-associated antigenic target (TAT) cDNA DNA327102, SEQ ID NO:6302.
W02004030615-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.7%; Score 67.2; DB 10; Length 2547; 90.0%; Pred. No. 2.9e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.7%; Score 67.2; DB 11; Length 2647; 90.0%; Pred. No. 2.9e-07;
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                                                                                                                                                                                                                                              ) 15-APR-2004.
(GETH) GENENTECH INC.
Query Match
Best Local Similarity 90.0%; Pred. No. 2.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HELI-) HELIX RES INST.
ry Match
2.7%; Score 67.2; DB 4; Length 2675; E Local Similarity 90.0%; Pred. No. 3e-07;
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(INCY-) INCYTE PHARM INC.

2.7%; Score 67.2; DB 2; Length 2713; ery Match

2.7%; Pred. No. 3e-07;
2.7%; Score 67.2; DB 5; Length 2947; 90.0%; Pred. No. 3.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX7233 standard; DNA; 2713 BP.
Human Rab protein, RABP-3, coding sequence.
WO9909182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADM01722 standard; cDNa; 2647 BP.
Human cDNa of the invention SEQ ID NO:407.
E14-SEP-2003.
(REAS-) RES ASSOC BIOTECHNOLOGY.
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Rat cDNA encoding tricarboxylate carrier
WO200138369-A1.
                                                                                                                                                                                                                                                                                                                                                      ADB62665 standard; cDNA; 2547 BP.
Human cDNA encoding clone LIVER20010510.
EP1308459-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human PRO cDNA sequence SEQ ID NO:2198 WO2004039956-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAH17304 standard; cDNA; 2675 BP.
Human cDNA sequence SEQ ID NO:16707.
EP1074617-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADNO5948 standard; cDNA; 2894 BP. Antipsoriatic cDNA sequence #1208 WO2004028479-A2.
                                                                                                                                                                              ACN41092 standard; cDNA; 2407 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADP56222 standard; cDNA; 2894 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                             (HELI-) HELIX RES INST.
(REAS-) RES ASSOC BIOTECHNOLOGY.
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(BIOR-) BIOROAD GENE DEV LTD 2.7%;
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(GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-APR-2004.
(GETH ) GENENTECH INC.
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(REAS-) RES ASSOC BIOTECHNOLOGY.

17%; Score 67.2; DB 12;

1 Local Similarity 90.0%; Pred. No. 3.2e-07;
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(LIFE-) LIFESPAN BIOSCIENCES INC.
                                                                                                                                                       (REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HUMA-) HUMAN GENOME SCI INC.
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                                     Query Match
Best Local Similarity
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28-JUL-2004
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              AAK79326 standard; DNA; 2997 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:34138.
WO200157183-A2.
09-AUG-2001.
(HUMA.) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 4; Length 2997;
st Local Similarity 90.0%; Pred. No. 3.1e-07;
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PA (HUMA-) HUMAN GENOME SCI INC.

Query Match

2.7%; Score 67.2; DB 10; Length 2997;

RESEL Local Similarity 90.0%; Pred. No. 3.1e-07;

RESULT 793
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                                                                                                                                                                                             Genomic sequence #35 encoding novel human DNA-binding protein. WO200155162-A1. 02-AUG-2001.
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Pred. No. 3.2e-07;
                                                                                                                                                                                                                                                                              Length 2997
                                                                                                                                                                                                                                                                                                                                  ABS68332 standard; DNA; 2997 BP.
Human DNA-binding protein genomic DNA sequence #35.
US2002102638-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADK60442 standard; DNA; 3139 BP.
Angiogenesis differentially expressed gene GS-N16.
FR2836687-A1.
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Angiogenesis differentially expressed gene GS-N16.
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Angiogenesis inhibitor human DNA sequence, GS-N16.
FR2843753-A1.
                                                                                                                                                                                                                                                                            2.7%; Score 67.2; DB 5; 90.0%; Pred. No. 3.1e-07;
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90.0%; Pred. No. 3.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADC25454 standard; cDNA; 2997 BP.
Human cDNA from extracellular matrix gene 60 #2.
US2003049650-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Signal pepiidase 11.88 cDNA.
CN1352292-A.
05-JUN-2002.
(BODE-) BODE GENE DEV CO LTD SHANGHAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADQ63046 standard; cDNA; 3151 BP.
Novel human cDNA sequence #207.
EP1440981-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABV72847 standard; cDNA; 3078 BP
                                                                                                                                                                          AAS29192 standard; DNA; 2997 BP
                                                                                                                                                                                                                                                       (HUMA-) HUMAN GENOME SCI INC.
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05-SEP-2003.
(GENE-) GENE SIGNAL.
(ALMA/) AL MAHMOOD S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GENE-) GENE SIGNAL.
(ALMA/) AL MAHMOOD S.
                                                                                                                              Best Local Similarity RESULT 790
                                                                                                                                                                                                                                                                                                                                                                                                                 (ROSE/) ROSEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                               Local Similarity
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Best Local Si
RESULT 796
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RESULT 794
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ADROBIIO standard; cDNA; 3268 BP.
Full length human cDNA useful for treating neurological disease Seq 1616.
EP1447413-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADR07160 standard; cDNA; 3494 BP. Full length human cDNA useful for treating neurological disease Seq 666. EP1447413-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAK75977 standard; DNA, 3452 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:30789.
WO200157182-A2.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:20385.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAL07337 standard; DNA; 4059 BP.
Human reproductive system related antigen DNA SEQ ID NO: 10025
WO200155320-A2.
                                                                                                                                                                                                                               2.7%; Score 67.2; DB 13; Length 3268; 90.0%; Pred. No. 3.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 3494;
Length 3151
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Human C-C chemokine receptor 6 nucleotide SBQ ID NO:389
WO200261087-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PD 18-AUG-2004.

PA (REAS-) RES ASSOC BIOTECHNOLOGY.

Query Match
2.7%; Score 67.2; DB 13;
Best Local Similarity 90.0%; Pred. No. 3.4e-07;
RESULT 802
                                                                                                                                                                                                                                                                                                                                                                                                                              (MCLA-) MCLAUGHLIN RES INST.
ry Match
t Local Similarity 90.0%; Pred. No. 3.4e-07;
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2.7%; Score 67.2; DB 4;

t Local Similarity 90.0%; Pred. No. 3.7e-07;
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2.7%; Score 67.2; DB 6;
Local Similarity 90.0%; Pred. No. 4.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABV77901 standard; DNA; 5779 BP. Hypoxia-repressed protein coding sequence #12.W0200246465-A2.
                                                                                                                                                                                                                                                                                                                ABZ80230 standard, cDNA, 3434 BP.
Human tramdorin 1 encoding cDNA SEQ ID NO:3.
WO2003016502-A2.
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Ouery Match Best Local Si RESULT 808

Query Match

Quer,
Best Loc.
RESULT 809
ID ABZ09P
DE Hummar
PN WO?
PD 0'

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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:37920.
WO200157182-A2.
09-AUG-2001.
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Human testicular antigen encoding DNA fragment SEQ ID NO: 2813
WO200155317-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.7%; Score 67.2; DB 6; Length 10139; 90.0%; Pred. No. 6.2e-07;
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Genomic sequence #69 encoding novel human DNA-binding protein.
WO200155162-A1.
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Human reproductive system related antigen DNA SEQ ID NO: 7965.
WO200155320-A2.
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2.7%; Score 67.2; DB 4; Length 8253; 90.0%; Pred. No. 5.5e-07;
                                                                                                                                                                                                                                              Length 8253;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABAI6439 standard; DNA; 8994 BP.
Human nervous system related polynucleotide SEQ ID NO 8770.
WO200159063-A2.
                                                                                                                                                                                                                                                                                                                                                           Human prostate cancer associated gene-related DNA SeqID622. US2003054373-A1.
                                                                                                                         prostate cancer antigen, Seq ID No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human DNA-binding protein genomic DNA sequence #69
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2.7%; Score 67.2; DB 11;

t Local Similarity 90.0%; Pred. No. 5.5e-07;
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(HUMA-) HUMAN GENOME SCI INC.

2.7%; Score 67.2; DB 5;

Lery Match 2.7%; Pred. No. 5.8e-07;
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90.0%; Pred. No. 6.4e-07;
                                                                                                                                                                                                                                              2.7%; Score 67.2; DB 5; 90.0%; Pred. No. 5.5e-07;
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Human cDNA from extracellular matrix gene 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAK83108 standard; DNA; 10914 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABS68366 standard; DNA; 10139 BP
                                                                                                                                                                                                                                                                                                                                           ADJ09676 standard; DNA; 8253 BP.
                                                                                               DNA; 8253 BP
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(HUMA-) HUMAN GENOME SCI INC.
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(ROSE/) ROSEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
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Best Local Similarity
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RESULT 821
                               Best Local Similarity RESULT 816
                                                                                         AAS40470 standard;
DNA encoding human
WO200155316-A2.
                                                                                                                                                                                                                                                                           Local Similarity
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RESULT 823
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Human cardiovascular system associated polypeptide-related DNA SeqID2327.
US2004005575-A1.
08-JAN-2004.
(HUMA-) HUMAN GENOME SCIINC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human cardiovascular system antigen genomic DNA SEQ ID No 2327 WO200155321-A2.
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Human reproductive system related antigen DNA SEQ ID NO: 6775.
WO200155320-A2.
                                      2.7%; Score 67.2; DB 12; Length 5779; 90.0%; Pred. No. 4.5e-07;
                                                                                                                                                                                                                                                                                     DB 13; Length 5779;
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                                                                                                                                                                                                                                                                                                                                                                    ABZ09932 standard; DNA; 6621 BP.
Human 5' and/or regulatory region of CDC25A DNA SEQ ID NO:72.
WO200277272-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABZ09875 standard; DNA; 6621 BP.
Human 5' and/or regulatory region of CDC25A DNA SEQ ID NO:15.
WO200277272-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABZ67712 standard; DNA; 7960 BP.
Human secreted protein encoding genomic DNA SEQ ID NO 1235.
020027186-A2.
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US2001059908-A1.
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(HUMA-) HUMAN GENOME SCI INC.

ery Match
2.7%; Score 67.2; DB 10;

ery Match
3.7%; Pred. No. 5.4e-07;
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02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 4;
ery Match 2.7%; Score 67.2; DB 4;
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Pred. No. 4.9e-07;
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Pred. No. 4.9e-07;
                                                                                                                                                                                                                        21-MAY-2004.
(GETH) GENENTECH INC.
2.7%; Score 67.2; DB 13
st Local Similarity 90.0%; Pred. No. 4.5e-07;
                                                                                                                            ADP22971 standard; cDNA; 5779 BP.
PRO polypeptide encoding cDNA SEQ ID NO:65.
WO2004041170-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADE84011 standard; DNA; 6621 BP.
5' regulatory region of human gene CDC25A.
WO2003044226-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAS36827 standard; DNA; 7960 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADE47521 standard; DNA; 7960 BP
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03-007-2002.
(EPIG-) EPIGENOMICS AG.
2.7%;
match ''nrity 90.0%;
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(EPIG-) EPIGENOMICS AG.
2ry Match 2.7%;
ELocal Similarity 90.0%;
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(EPIG-) EPIGENOMICS AG.
   PA (GETH ) GENENTECH INC.
Query Match
Best Local Similarity 9
RESULT 807
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Best Local Similarity
RESULT 814
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Query Match

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Best Loca RESULT 810

Query Match

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us-10-002-967a-58.rng.spdi

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y Match 2.7%; Score 67.2; DB 12; Length 12565; Local Similarity 90.0%; Pred. No. 7e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADL81319 standard; DNA; 12565 BP.
Human secreted interleukin-1 receptor antagonist (IL-1RN) gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 12565;
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  2.7%; Score 67.2; DB 5; Length 12565; 90.0%; Pred. No. 7e-07;
                                                                                                                                                                            Length 12565;
                                                                   Human interleukin 1 receptor antagonist genomic sequence WO2003026488-A2.
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Pred. No. 7e-07;
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Local Similarity 90.0%; Pred. No. 7e-07;
                                                                                                                                                                         Score 67.2; DB 8;
Pred. No. 7e-07;
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                                                                                                                                                                                                                                       AAL54517 standard; DNA; 12565 BP.
Secreted interleukin-1RN nucleotide sequence.
WO2002103031-A2.
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WO2004041170-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADN48840 standard; DNA; 12565 BP.
Human secreted interleukin-1RN (IL-1RN) DNA.
US6730476-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADO24452 standard; cDNA; 12565 BP.
Human DNA327777 cDNA clone SEQ ID NO:91
                                                                                                                                                                                                                                                                                                                                                                                                                   AAD51465 standard, DNA, 12565 BP.
Human interleukin-1RN (IL-1RN) gene.
WO2002101015-A2.
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Human interleukin-1RN (IL-1RN) gene.
US6746839-B1.
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                                                                                                                                                                                                                                                                                                                                 INC.
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(INTE-) INTERLEUKIN GENETICS INC.
ry Match
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(INTE-) INTERLEUKIN GENETICS INC.
                                                                 ACC44353 standard; DNA; 12565 BP.
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(INTE-) INTERLEUKIN GENETICS INC.
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WO2004028479-A2.
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(INTE-) INTERLEUKIN GENETICS INC.
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PD 27-MAY-2004.
PA (GETH) GENENTECH INC.
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                                                                                                                                                       (VITI-) VITIVITY INC.
                  Best Local Similarity RESULT 834
                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
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Local Similarity
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                                                                                                                                                                                                                                                           D AAK81675 standard; DNA; 11819 BP.

E Human immune/haematopoietic antigen genomic sequence SEQ ID NO:36487.

N WO200157182-A2.

D 99-AUG-2001.

A (HUMA-) HUMAN GENOME SCI INC.

Query Match

2.7%; Score 67.2; DB 4; Length 11819;

Best Local Similarity 90.0%; Pred. No. 6.7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human immune/haematopoietic antigen genomic sequence SEQ ID NO:36722.
WO200157182-A2.
                        2.7%; Score 67.2; DB 4; Length 10914; 90.0%; Pred. No. 6.4e-07;
                                                                                                                                                                                                Length 11173;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PD 09-AUG-2001.

PA (HUMA-) HUMAN GENOME SCI INC.

Query Match 2.7%; Score 67.2; DB 4; Length 12466;

Best Local Similarity 90.0%; Pred. No. 6.9e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-001-1998.
(GEMI-) GEMINI RES LTD.
ery Match
ery Match
ery Match
2.7%; Score 67.2; DB 2; Length 12565;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 12565;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.7%; Score 67.2; DB 3; Length 12565; 90.0%; Pred. No. 7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.7%; Score 67.2; DB 4; Length 12565; 90.0%; Pred. No. 7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 2; Length 12565;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human interleukin-1 receptor antagonist, IL-IRN, US2002146700-A1.
                                                                                                                                                   PD 16-JAN-2003.

PA (1S.S-) ISIS PHARM INC.

Query Match 2.7%; Score 67.2; DB 9;

Best Local Similarity 90.0%; Pred. No. 6.5e-07;

RESULT 826
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDI-) MEDICAL SCI SYSTEMS INC.
(MEDI-) MEDICAL SCI SYSTEMS INC.
2.7%; SCOIC 67.2; DB 2
ery Match
Cimilarity 90.0%; Pred. No. 7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OCCT-2002.

(INTE-) INTERLEUKIN GENETICS INC.

ery Match
2.7%; Score 67.2; DB 2
ery Match
7e-7 Similarity 90.0%; Pred. No. 7e-07;
                                                                                      ABZ70982 standard; DNA; 11173 BP.
Human HKR1 related DNA sequence SEQ ID NO:10.
WO2003004513-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human interleukin-1 receptor antagonist gene.
WO9844150-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAC63769 standard; DNA; 12565 BP.
Human IL-1RN gene.
WO200060117-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (INTE-) INTERLEUKIN GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-OCT-2000.
(INTE-) INTERLEUKIN GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA; 12565 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200100880-A2.
04-JAN-2001.
(INTE-) INTERLEUKIN GENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAC91435 standard; DNA; 12565 BP.
Human IL-1RN nucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA; 12565 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA; 12466 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV62392 standard; DNA; 12565 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABX15530 standard; DNA; 12565 BP
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAX75925 standard; DNA; 125
Human interleukin 1RN gene.
WO9924615-A2.
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                                      Best Local Similarity RESULT 825
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
RESULT 829
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
RESULT 831
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                            AAK81910 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF27667 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200071753-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IL-1RN DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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RESULT 830

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AAK70808 standard; DNA; 15484 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:25620.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAS36826 standard; DNA; 17239 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADE47520 standard; DNA; 17239 BP
                                                                                                                                                                                                                                                                                               2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity RESULT 858
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
RESULT 861
                                                                                                                         Query Match
Best Local Similarity
RESULT 853
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Best Local Similarity
                                                                                                                                                                                                                                                       25-FEB-1999.
(FARB ) BAYER CORP.
                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                               WO9909150-A1.
25-FEB-1999.
(FARB ) BAYER CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity RESULT 855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (FARB ) BAYER CORP. Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                         AAX24732 standard;
Swedish/London-FAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-FEB-1999
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      RESULT 852
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ID AAK73722 standard; DNA; 14085 BP.

DE Human immune/haematopoietic antigen genomic sequence SEQ ID NO:28534.

PN W0200157182-A2.

PD 09-AUG-2001.

PA (HUMA-) HUMAN GENOME SCI INC.

QUETY Match

2.7%; Score 67.2; DB 4; Length 14085;

BB81 Local Similarity 90.0%; Pred. No. 7.49-07;
                                                                                                                                                                                           AAZ32189 standard; DNA; 13204 BP.
Human platelet glycoprotein IIb exons 2-29 nucleotide sequence.
WO9950454-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 13392;
                                                                                                                                                DB 2; Length 12814;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      A (HUMA-) HUMAN GENOME SCI INC.
Query Match
2.7%; Score 67.2; DB 4; Length 13392;
Best Local Similarity 90.0%; Pred. No. 7.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 15186;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 13392;
                                                                                                                                                                                                                                                                                                                  Length 13204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 14231
                                                                                                                                                                                                                                                                                                                                                                                             Human reproductive system related antigen DNA SEQ ID NO: 6774 WO200155320-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABA19751 standard; DNA; 14231 BP.
Human nervous system related polynucleotide SEQ ID NO 12082
WO200159063-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABA15617 standard; DNA; 15186 BP.
Human nervous system related polynucleotide SEQ ID NO 7948.
WO200159063-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABAIS616 standard; DNA; 15187 BP.
Human nervous system related polynucleotide SEQ ID NO 7947.
WQ200159063-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADJ09675 standard; DNA; 13392 BP.
Human prostate cancer associated gene-related DNA SeqID621
US2003054373-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAS40469 standard; DNA; 13392 BP.
DNA encoding human prostate cancer antigen, Seq ID No 621.
WO200155316-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PD 20-MAR-2003.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 2.7%; Score 67.2; DB 11;
Best Local Similarity 90.0%; Pred. No. 7.2e-07;
RESULT 848
                                                                                                                                                                                                                                                                 07-OCT-1999.
(WHED ) WHITEHEAD INST BIOMEDICAL RES.
2.7%; Score 67.2; DB 2;
cery Match 2.7%; Pred. No. 7.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PD 16-AUG-2001.

PA (HUMA-) HUMAN GENOME SCI INC.

QUERY Match
2.7%; Score 67.2; DB 5;
Best Local Similarity 90.0%; Pred. No. 7.7e-07;

RESULT 851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 5;
it Local Similarity 90.0%; Pred. No. 7.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.7%; Score 67.2; DB 5; 90.0%; Pred. No. 7.5e-07;
                                                                                                                                              2.7%; Score 67.2; DB : 90.0%; Pred. No. 7e-07;
    90.0%; Pred. No. 7e-07;
                                        ID AAX24730 standard; DNA; 12814 BP.
DE Swedish-FAD APP targetting vector pWTI-2398.
PN W09909150-A1.
PD 75-FEB-1999.
PA (FARB ) BAYER CORP.
                                                                                                                                                                                                                                                                                                                                                                          AAL04086 standard; DNA; 13392 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                           Best Local Similarity
RESULT 844
ID AA232189 standard; Di
DE Human platelet glycof
PN W09950454-42.
PD 07-00T-1999.
PA (WHED ) WHITEHEAD INS
Best Local Similarity RESULT 843
                                                                                                                                                                                                                                                                                                                               Best Local Similarity RESULT 845
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Best Local Similarity
RESULT 850
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                       02-AUG-200
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Best Local Si
RESULT 847
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AAK79798 standard; DNA; 16682 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:34610.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAK70010 standard; DNA; 16682 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:24822.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-OCT-2002.
(HUMA-) HUMAN GENOME SCI INC.
(ery Match 2.7%; Score 67.2; DB 10; Length 17239;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human cardiovascular system antigen genomic DNA SEQ ID No 2326.
WO200155321-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 17239;
                                                                                                                                                                                                                                                               Length 15692;
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                                                                Length 15484;
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Human secreted protein encoding genomic DNA SEQ ID NO 1234
WQ200277186-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human cardiovascular system related genomic DNA #1086
US2003059908-A1.
                                                                                                                                                                                                                                                                                                                                          DNA; 15692 BP.
APP targetting vector pMTI-2454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.7%; Score 67.2; DB 10; 90.0%; Pred. No. 8.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.7%; Score 67.2; DB 2; 90.0%; Pred. No. 7.9e-07;
MOLOCOLOGO

09-MUCA-) HUMAN GENOME SCI INC.

(HUMA-) HUMAN GENOME SCI INC.

2.7%; Score 67.2; DB 4;

lery Match

2.7%; Score 67.2; DB 4;

2.7%; Pred. No. 7.8e-07;
                                                                                                                                                                                                                                                                 Score 67.2; DB 2;
Pred. No. 7.9e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WOLVELL.
02-AUG-2001.
(HUMAL) HUMAN GENOME SCI INC.
(HUMAL) AUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 4;
tery Match
2.7%; Pred. No. 8.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 67.2; DB 2;
Pred. No. 7.9e-07;
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(HUWA-) HUMAN GENOME SCI INC.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 4;
ery Match
2.7%; Pred. No. 8.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HUMA-) HUMAN GENOME SCI.INC.

2.7%; Score 67.2; DB 4;

L Local Similarity 90.0%; Pred. No. 8.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAXZ-1733 standard; DNA; 15701 BP.
Swedish-FAD APP713 targetting vector pMTI-2454.
WO9909150-A1.
                                                                                                                                   AAX24731 standard; DNA; 15692 BP.
London-FAD APP targetting vector pMTI-2453
WO9909150-A1.
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Length 20245;

Length 20245

Length 20245;

Length 20245;

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AAK72318 standard; DNA; 20245 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:27130.
WO200157182-A2.
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WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABZ71482 standard; DNA; 20245 BP.
Secreted protein gene 12 genomic fragment HAUAI83, SEQ ID NO:592
WO200276488-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.7%; Score 67.2; DB 10; Length 20245; 90.0%; Pred. No. 9.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.7%; Score 67.2; DB 10; Length 20245; 90.0%; Pred. No. 9.1e-07;
  HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 4; Length 20068;
Similarity 90.0%; Pred. No. 9e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADAS7727 standard; DNA; 20245 BP.
BAC fragment containing human secreted protein gene #302 WO2002102994-A2.
                                                                                                                                                                                                                                                                        ABK69845 standard; DNA; 20245 BP.
Human secreted protein gene 22 genomic DNA fragment #6
WO200226931-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.7%; Score 67.2; DB 10; 90.0%; Pred. No. 9.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADC74613 standard; DNA; 20245 BP.
Human secreted protein-related DNA - SEQ ID 1246.
                                                                                                                                                                   09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 4;
EL Local Similarity 90.0%; Pred. No. 9.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          03-OCT-2002.
(HUMA-) HUMAN GENOME SCI INC.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 8;
ery Match
2.7%; Pred. No. 9.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                     2.7%; Score 67.2; DB 6; 90.0%; Pred. No. 9.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACCS0860 standard; cDNA; 20245 BP.
Human secreted protein BAC clone SEQ ID NO 1040.
WO2002.9010.A2.
28-NOV-2002.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HUMA-) HUMAN GENOME SCI INC.
ry Match
t Local Similarity 90.0%; Pred. No. 9.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.7%; Score 67.2; DB 8; 90.0%; Pred. No. 9.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human secreted protein related DNA #SEQ ID 784. W0200304622-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CDNA; 20245 BP. deposit #27.
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                                                                                                                                                                                                                                                                                                                                                                           (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-MAY-2003.
(HUMA-) HUMAN GENOME SCI INC.
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(HUMA-) HUMAN GENOME SCI INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity RESULT 873
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
RESULT 874
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                            Query Match
Best Local Similarity
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877
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADD38133 standard;
cDNA clone in ATCC
WO200290526-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-JAN-2003
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                                                                                                                                                                                                                 Query Match
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       (HUMA-)
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ADJ08938 standard; DNA; 17239 BP.

Human cardiovascular system associated polypeptide-related DNA SeqID2326.
US2004005575-A1.
08-JAN-2004.
(HUMA-N FENOME SCI INC.
2.7%; Score 67.2; DB 13; Length 17239;
ELocal Similarity 90.0%; Pred. No. 8.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAK70445 standard; DNA; 18949 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:25257.
WO200157182-A2.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:21545.
WO200157182-A2.
09-AUG-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAK87416 standard; DNA; 19942 BP.
Huwan immune/haematopoietic antigen genomic sequence SEQ ID NO:42228.
WO200157182-A2.
09-AUG-2001.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:21547.
WO200157182-A2.
                                                                                                                                                                 AAL136869 standard, DNA, 17286 BP.
Human musculoskeletal system related polymucleotide SRQ ID NO 3234.
WO200155367-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADJ30607 standard; DNA; 17286 BP.
Human musculoskeletal system-associated genomic DNA - SEQ ID 3234
US2004009488-A1.
                                                                                                                                                                                                                                                                                                                                                        ABX59857 standard; cDNA; 17286 BP.
CDNA encoding novel human musculoskeletal system antigen #2201.
US2002147140-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.7%; Score 67.2; DB 12; Length 19913; 90.0%; Pred. No. 9e-07;
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(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 4; Length 18949;
ery Match
2.7%; Score 67.2; DB 4; Length 18949;
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cy Match 2.7%; Score 67.2; DB 4; Length 20067;
c Local Similarity 90.0%; Pred. No. 9e-07;
                                                                                                                                                                                                                                                                                                            Length 17286
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(HUWA-) HUMAN GENOME SCI INC.
ry Match
t Local Similarity 90.0%; Pred. No. 8.3e-07;
                                                                                                                                                                                                                                                                             (HUMA-) HUMAN GENOME SCI INC.

2.7%; Score 67.2; DB 4;
t Local Similarity 90.0%; Pred. No. 8.3e-07;
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MSI-H carcinoma genomic DNA sequence SEQ ID NO:2.
KR2004008012-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.7%; Score 67.2; DB 8; 90.0%; Pred. No. 8.3e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ROSE/) ROSEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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(KIMN/) KIM N G.
(LEEJ/) LEE J S.
(RHEE/) RHEE H S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-JAN-2004.
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Best | RESULT (

Length 20245;

Length 20245;

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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:22456.
WO200157182-A2.
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Secreted protein gene 310 genomic fragment HSYBI06, SEQ ID NO:1626.
WO200277013-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       / Match 2.7%; Score 67.2; DB 13; Length 24810; Local Similarity 90.0%; Pred. No. 1e-06;
                     2.7%; Score 67.2; DB 10; Length 21784; 90.0%; Pred. No. 9.4e-07;
                                                                                                                                                                                                                                                                                   2.7%; Score 67.2; DB 10; Length 21784; 90.0%; Pred. No. 9.4e-07;
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Human testicular antigen encoding DNA fragment SEQ ID NO: 3045.
WO200155317-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.7%; Score 67.2; DB 4; Length 23544; 90.0%; Pred. No. 9.9e-07;
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Human reproductive system related antigen DNA SEQ ID NO: 8517.
WO200155320-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 21982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABZ68026 standard; DNA; 21982 BP.
Human secreted protein encoding genomic DNA SEQ ID NO 1549
WO200277186-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADQ97272 standard; DNA; 24861 BP.
Human cancer associated sequence HD08-024, SEQ ID 248.
WO2004060304-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WACCOLL
21-NOV-2002.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 10;
ery Match
2.7%; Score 67.2; DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADC20924 standard; DNA; 21982 BP.
Human secreted protein-related DNA sequence #342.
WO200292787-A2.
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03-OCT-2002.
(HUMA-) HUMAN GENOME SCI INC.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 8;
ery Match
2.7%; Pred. No. 9.5e-07;
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2.7%; Score 67.2; DB 4;
t Local Similarity 90.0%; Pred. No. 9.9e-07;
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Human cancer-associated genomic DNA HD13-026.
WO2004074320-A2.
                                                                                     ABX16008 standard; DNA; 21784 BP.
Human novel protease gene.
US2002142440.Al.
03-OCT-2002.
(CANW/) GAN W.
(VEJJ/) YB J.
(BFAA/) BEASLEY E M.
                                                                                                                                                                                                                                                                                                                            RESULT 889
ID AAK67644 standard; DNA; 21982 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                        09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
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(HUMA-) HUMAN GENOME SCI INC.
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(SAGR-) SAGRES DISCOVERY INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
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Best Local Similarity
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Best Local Similarity
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Best Local Similarity
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                                           Best Local Similarity RESULT 888
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Best Local Si
RESULT 896
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RESULT 894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human cardiovascular system associated polypeptide-related DNA SeqID1734. US2004005575-A1.
                                                       ID AAK79760 standard, DNA; 20601 BP.

DB Human immune/haematopoietic antigen genomic sequence SEQ ID NO:34572.

PD 9-AUG-2001.

PA (HUMA-) HUMAN GENOME SCI INC.

Query Match
2.7%; Score 67.2; DB 4; Length 20601;

Best Local Similarity 90.0%; Pred. No. 9.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      _AAK79799 standard; DNA; 21436 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:34611.
WO200157182-A2.
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WO200157182-A2.
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DE Human GPCR gene SEG ...

PN EP1270724-A2.

PN 02-JAN-2003.

PA (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.

PA (ADSC-) CENT ADVANCED SCI & TECHNOLOGY INCUBATIO.

Query Match

2.7%; Score 67.2; DB 10; Length 21581;

Best Local Similarity 90.0%; Pred. No. 9.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-MAR-2003.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 10; Length 21129;
ELocal Similarity 90.0%; Pred. No. 9.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PD 08-JAN-2004.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 2.7%; Score 67.2; DB 13; Length 21129;
Best Local Similarity 90.0%; Pred. No. 9.3e-07;
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PA (HUMA-) HUMAN GENOME SCI INC.

Query Match

2.7%; Score 67.2; DB 4; Length 21436;

Best Local Similarity 90.0%; Pred. No. 9.4e-07;

RESULT 886
                                                                                                                                                                                                                                                                                AAS36234 standard; DNA; 21129 BP.
Human cardiovascular system antigen genomic DNA SEQ ID No 1734
WO200155321-A2.
                                                                                                                                                                                                                                                                                                                                               02-Aug-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match imilarity 90.0%; Pred. No. 9.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                uuery Match 2.7%; Score 67.2; DB 6; Length 21234;
Best Local Similarity 90.0%; Pred. No. 9.3e-07;
RESULT 884
Query Match 2.7%; Score 67.2; DB 4; Length 20323; Best Local Similarity 90.0%; Pred. No. 9.1e-07; RESULT 879
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 21436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADE46928 standard; DNA; 21129 BP.
Human cardiovascular system related genomic DNA #494
US2003059908-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
(ery Match 2.7%; Score 67.2; DB 4;
ery Match 90.0%; Pred. No. 9.4e-07;
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ID AAK70011 standard; DNA; 21436 BP. DE Human immune/haematopoietic antigen PN WO200157182-A2. PD 09-AUG-2001. PA (HUMA-) HUMAN GENOME SCT INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAD32039 standard; DNA; 21234 BP. Human kinase protein gene. 20200216567-A2. 28-FEB-2002. (APPL-) APPLERA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA; 21129 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-JUL-2003.
(APPL-) APPLERA CORP.
                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
RESULT 881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADJ08346 standard;
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Best Local Si
RESULT 882
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Length 21982;

Query Match

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AAK81578 standard; DNA, 32313 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:36390.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAK67071 standard; DNA; 33923 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:21883.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAK73093 standard; DNA; 33747 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:27905.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAK69279 standard; DNA; 33747 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:24091
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.7%; Score 67.2; DB 13; Length 34796; 90.0%; Pred. No. 1.2e-06;
                                  02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 67.2; DB 4; Length 31931;
-- ' ... Similarity 90.0%; Pred. No. 1.2e-06;
                                                                                                                                                                  Human reproductive system related antigen DNA SEQ ID NO: 9663 WO200155320-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match
ery Match
-- '- '- ' Similarity 90.0%; Pred. No. 1.2e-06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACN37240 standard; DNA; 34796 BP.
Human periodontal disease related gene PLOD SEQ ID NO:150.
WO2004042054-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human soft tissue sarcoma-upregulated DNA - SEQ ID 1073 WO2004048938-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (PROT-) PROTEIN DESIGN LABS INC.

(PROT-) PROTEIN DESIGN LABS INC.

2.7%; Score 67.2; DB 12;

ery Match

2.7%; Pred. No. 1.2e-06;
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PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 2.7%; Score 67.2; DB 4;
Best Local Similarity 90.0%; Pred. No. 1.2e-06;
RESULT 908
                                                                                                                                                                                                        02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 67.2; DB 4;
ery Match 90.0%; Pred. No. 1.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 67.2; DB 4;
Pred. No. 1.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.7%; Score 67.2; DB 6; 90.0%; Pred. No. 1.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HUMAN GENOME SCI INC.
1 2.7%; Score
Similarity 90.0%; Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADC87616 standard, DNA, 35425 BP.
Human GPCR gene SEQ ID NO:2069.
EP1270724-A2.
                                                                                                                                                  DNA; 32250 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA; 33414 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAD44328 standard; DNA; 34667 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HUBI-) HUBIT GENOMIX INC.
(KAMO/) KAMOI K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human transporter gene. WO200248367-A2.
                                                                                               Best Local Similarity RESULT 906
                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity
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                                                                                                                                             AAL06975 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADQ18255 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200157182-A2.
                  WO200155206-A1.
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                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABA08089 standard; DNA; 31304 BP.
Human ovarian and breast cancer associated polynucleotide SEQ ID NO 884.
WO200155325-A2.
                                                                                                                                                                                                                                                                                  AAK67239 standard, DNA, 30393 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:22051.
W0200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AALJ7445 standard; DNA; 31730 BP.
Human musculoskeletal system related polynucleotide SEQ ID NO 3810.
WO200155367-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADJ31183 standard; DNA; 31730 BP.
Human musculoskeletal system-associated genomic DNA - SEQ ID 3810
US2004009488-A1.
PA (SAGR-) SAGRES DISCOVERY INC.
QUERY MArch
Best Local Similarity 90.0%; Pred. No. 1e-06;
                                                                                                                                                                                          12-SEP-2003.
(SAGR.) SAGRES DISCOVERY.
2.7%; Score 67.2; DB 11; Length 30030;
it Local Similarity 90.0%; Pred. No. 1.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 67.2; DB 11; Length 31652; Pred. No. 1.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CDNA encoding novel human musculoskeletal system antigen #2777.
US2002147140-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human reproductive system related antigen DNA SEQ ID NO: 9556. WO200155320-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                       Length 30393
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Human pancreatic cancer related genomic DNA, SEQ ID NO: 479.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.7%; Score 67.2; DB 12; 90.0%; Pred. No. 1.2e-06;
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(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 67.2; DB 4;
ery Match 90.0%; Pred. No. 1.2e-06;
                                                                                                                                                                                                                                                                                                                                                               09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
(ELY Match 2.7%; Score 67.2; DB 4;
ery Match 90.0%; Pred. No. 1.1e-06;
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(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 67.2; DB 4;
ery Match 2.7%; Score 67.2; DB 4;
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                                                                                                                      ACN44714 standard; DNA; 30030 BP.
Human genomic sequence hCG14997.
WO2003073826-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACN44118 standard; DNA; 31652 BP.
Human genomic sequence hCG23908.
WO2003073826-A2.
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(HUMA-) HUMAN GENOME SCI INC.
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Local Similarity 90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (SAGR-) SAGRES DISCOVERY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RUBE/) RUBEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAL06868 standard;
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Best Local Si
RESULT 900
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Best RESULT

Length 34667;

Query Match

Length 33923;

Match

Length 33414;

Length 32250;

Length 32313;

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ABN97975 standard; DNA; 44100 BP
           WO2003073826-A2.
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RESULT
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DE Secreted protein gene 299 genomic fragment HSOAH16, SEQ ID NO:1606.
PN WO200277013-A2.
PD 03-OCT-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match
2.7%; Score 67.2; DB 8; Length 39339;
Best Local Similarity 90.0%; Pred. No. 1.3e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PD 03-OCT-2002.

PA (HUMA-) HUMAN GENOME SCI INC.

Query Match

Best Local Similarity 90.0%; Pred. No. 1.3e-06;

RESULT 923

ID ACM44798 standard; DNA: 4.4.7.
PA (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.

PA (ADSC-) CENT ADVANCED SCI & TECHNOLOGY INCHBATIO.

Query Match 2.7%; Score 67.2; DB 10; Length 35425;

Best Local Similarity 90.0%; Pred. No. 1.2e-06;
                                                                                                                                                                                          PA (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.

PA (ADSC-) CENT ADVANCED SCI & TECHNOLOGY INCHBATIO.

Query Match 2.7%; Score 67.2; DB 10; Length 35425;

Best Local Similarity 90.0%; Pred. No. 1.2e-06;
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DE Human lipase endothelial (LIPG) isogene genomic DNA sequence.

PN W0200216397-A2.

PD 28-FEB-2002.

PA (GENA-) GENAISSANCE PHARM INC.

Query Match 2.7%; Score 67.2; DB 6; Length 39328,

RESULT 921
                                                                                                                                                                                                                                                                                                                AAR'S4723 standard; DNA; 35465 BP.
Nucleotide sequence of a human polynucleotide sequence.
WO200105422-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 90.0%; Pred. No. 1.2e-06;
RESULT 918
ID ABD33508 standard; DNA; 36871 BP.
DE Human cancer-associated (CA) gene HD07-099.
PN W02004058146-A2.
PD 15-JUL-2004.
PA (SAGR-) SAGRES DISCOVERY INC.
Query Match
2.7%; Score 67.2; DB 13; Le.
RESULT 919
                                                                                                                                                                                                                                                                                                                                                                                     MCACLTO.
25-JAN-2001.
(IMR.) BIOMERIEUX STELHYS.
(Ery Match 2.7%; Score 67.2; DB 4;
ery Match 90.0%; Pred. No. 1.2e-06;
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Beet Local Similarity 90.0%; Pred. No. 1.3e-06;
RESULT 920
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Human IG gene related nucleic acid SEQ ID No
WO200299040-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABT17382 standard; DNA; 36991 BP.
Human IG gene related nucleic acid SEQ ID No
WO200299040-A2.
                                                                                                                 Human GPCR gene SEQ ID NO:2069.
EP1270724-A2.
02-JAN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-DEC-2002.
(EXEL-) EXELIXIS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-DEC-2002. (EXEL-) EXELIXIS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
RESULT 917
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ID ACM44166 standard; DNA; 41624 BP.

DE Human genomic sequence hCG1643838.

PN WO2003073826-A2.

PD 12-SEP-2003.

PA (SAGR-) SAGRES DISCOVERY.

Query Match 2.7%; Score 67.2; DB 11; Length 41624;

Best Local Similarity 90.0%; Pred. No. 1.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JUN-2003.

1 (SAGE) SAGRES DISCOVERY.

Query Match

Query Match

Best Local Similarity 90.0%; Pred. No. 1.4e-06;
                                       Query Match 2.7%; Score 67.2; DB 11; Length 40491; Best Local Similarity 90.0%; Pred. No. 1.3e-06;
                                                                                                                                                                                                                                                                                                           ACM45050 standard; DNA; 41966 BP.

Human genomic sequence hCG39344.
W02003073826-A2.
12-SEP-2003.
(SAGR-) SAGRES DISCOVERY.

ETY MAICH

EL Local Similarity 90.0%; Pred. No. 1.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.7%; Score 67.2; DB 11; Length 43936; 90.0%; Pred. No. 1.4e-06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        y Match 2.7%; Score 67.2; DB 10; Length 44030; Local Similarity 90.0%; Pred. No. 1.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABK84242 standard; cDNA; 43599 BP.
Human cDNA differentially expressed in granulocytic cells #813.
WO200228999-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 44030;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADA02906 standard; DNA; 44030 BP.
Human MCM3AP carcinoma associated gene, SEQ ID NO:1424
WO2003057146-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MCACCOLL
17-JUL 2003.
(SACH.) SAGRES DISCOVERY.
(SACH.) SAGRES DISCOVERY.
2.7%; Score 67.2; DB 9;
lery Match
2.7%; Pred. No. 1.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADM74501 standard; DNA; 44030 BP.
Human carcinoma associated (CA) nucleic acid #85
US2004072154-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-APR-2002.
(GENE-) GENE LOGIC INC.
Query Match
2.7%; Score 67.2; DB 6;
BEST Local Similarity 90.0%; Pred. No. 1.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADL95498 standard; DNA; 43936 BP. Human MCM3AP genomic DNA. US2003165878-A1. (MORK) MORRIS D W. (MORK) RNGELHARD E K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADB72644 standard; DNA; 44030 BP.
Human MCM3AP gene.
WO2003008583-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human Mcm3ap genomic sequence. WO2003045230-A2.
12-SEP-2003.
(SAGR-) SAGRES DISCOVERY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JAN-2003.
(SAGR-) SAGRES DISCOVERY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MORR/) MORRIS D W. (ENGE/) ENGELHARD E K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best_Local_Similarity
RESULT 928
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
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RESULT 947
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                                                                                                                             ABD32642 standard; DNA; 47096 BP.

ABD32642 standard; DNA; 47096 BP.

Human cancer-associated genomic DNA HD13-028.

W0204074320-A2.

W02-SEP-2004.

(SAGR-) SAGRES DISCOVERY INC.

2.7%; Score 67.2; DB 13; Length 47096; st Local Similarity 90.0%; Pred. No. 1.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 48001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 48469;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 67.2; DB 10; Length 47573;
Pred. No. 1.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 47573;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 47573;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 67.2; DB 11; Length 49536;
Pred. No. 1.5e-06;
                                                                            Length 44100;
                                                                                                                                                                                                                                                                                                                                                                                    Length 47573;
                                                                                                                                                                                                                                                                                    ADA02840 standard; DNA; 47573 BP.
Human MAP3KB carcinoma associated gene, SEQ ID NO:1358.
WO2003057146-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 67.2; DB 10;
Pred. No. 1.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC.
Query Match
2.7%; Score 67.2; DB 12;
Best Local Similarity 90.0%; Pred. No. 1.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Match 2.7%; Score 67.2; DB 11;
Local Similarity 90.0%; Pred. No. 1.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 2.7%; Score 67.2; DB 12;
Local Similarity 90.0%; Pred. No. 1.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human carcinoma associated (CA) nucleic acid #52 US2004072154-A1.
Human retroviral sequence RAM75.
W09967395-A1.
29-DEC-1999.
(INPM ) INSERM INST NAT SANTE & RECH MEDICALE.
ery Match
2.7%; Score 67.2; DB 3;
et Local Similarity 90.0%; Pred. No. 1.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                  Score 67.2; DB 9;
Pred. No. 1.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Low scandard; DNA; 49536 BP. Human genomic sequence hCG1775160. WOZ003073856-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACN44802 standard; DNA; 48469 BP.
Human genomic sequence hCG1640192
WO2003073826-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB72578 standard; DNA; 47573 BP.
Human MAP3K8 gene.
WO2003008583-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Abo61194 standard; DNA; 48001 BP. Human PPAR binding protein DNA. US2004101855-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADM74435 standard; DNA; 47573 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADC20666 standard; DNA; 51961 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA; 47573 BP.
                                                                                                                                                                                                                                                                                                                                   MOZOUREN
17-JUL-2003.
17-JUL-2003.
18AGR.) SAGRES DISCOVERY.
2.7%;
2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MCaco--
CACR-) SAGRES DISCOVERY.
(SAGR-) SAGRES DISCOVERY.
ery Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 90.0%; RESULT 936
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC85319 standard; DNA; 47573
Mouse Map3k8 coding sequence.
WO2003045230-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 2.7%;
Local Similarity 90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-SEP-2003.
(SAGR-) SAGRES DISCOVERY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-JAN-2003.
(SAGR-) SAGRES DISCOVERY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12-SEP-2003.
(SAGR-) SAGRES DISCOVERY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MORR/) MORRIS D W. (ENGE/) ENGELHARD E K.
                                                                                                                                                              Best Local Similarity RESULT 933
                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
RESULT 935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-APR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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15-MAY-2003.
(SAGR-) SAGRES DISCOVERY.
Query Match 2.7%; Score 67.2; DB 10; Length 52242;
Best Local Similarity 90.0%; Pred. No. 1.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                           03-OCT-2002.
(HUMA-) HUMAN GENOME SCI INC.
ery.Match 2.7%; Score 67.2; DB 10; Length 51961;
-- ' ' A Similarity 90.0%; Pred. No. 1.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 67.2; DB 10; Length 52242;
Pred. No. 1.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 67.2; DB 11; Length 58845;
Pred. No. 1.6e-06;
                                                                                                                                                                                                                 03-OCT-2002.
(HUMA-) HUMAN GENOME SCI INC.
ery Match
2.7%; Score 67.2; DB 10; Length 51961;
2.7%; Pred. No. 1.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 56577;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               / Match 2.7%; Score 67.2; DB 12; Length 57008; Local Similarity 90.0%; Pred. No. 1.6e-06;
                                                                                     Length 51961;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.7%; Score 67.2; DB 9; Length 52242; 90.0%; Pred. No. 1.5e-06;
                                                                                                                                                                    Human secreted protein-related DNA sequence - SEQ ID No 259 W0200277188-A2.
                                                                                                                                                                                                                                                                                                                          ABZ67488 standard; DNA; 51961 BP.
Human secreted protein encoding genomic DNA SEQ ID NO 1011.
WO200277186-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AUQ97617 standard; DNA; 57008 BP.
Human cancer associated sequence HD10-015, SEQ ID 594
WO2004060304-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human MDM2 carcinoma associated gene, SEQ ID NO:1184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human TK2 carcinoma associated gene, SEQ ID NO:1214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-58P-2003.
(SAGR-) SAGRES DISCOVERY.
2.7%; Score 67.2; DB 11;
rry Match 2.7%; Pred. No. 1.6e-06;
Human secreted protein-related DNA sequence #84. W0200292787-A2. 21-NOV-2002. (HUMA-) HUMAN GENOME SCI INC. 2.7%; Score 67.2; DB 10; St March 90.0%; Pred. No. 1.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADE95914 standard; DNA; 52242 BP.
Human MDM2 gene genomic DNA sequence.
WO2003039484-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACN43874 standard; DNA; 58845 BP.
Human genomic sequence hCG25637.
WO2003073826-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACN44778 standard; DNA; 56577 BP. Human genomic sequence hCG27352. WO2003073826-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADA02696 standard; DNA; 59554 BP
                                                                                                                                                    ABT16905 standard; DNA; 51961 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADA02666 standard; DNA; 52242 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADB72404 standard; DNA; 52242 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-JUL-2004.
(SAGR-) SAGRES DISCOVERY INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.7%;
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ry Match
t Local Similarity 90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO2003057146-A2.
17-JUL-2003.
(SAGR-) SAGRES DISCOVERY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (SAGR-) SAGRES DISCOVERY.
                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                      Query.Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
RESULT 945
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOZ003008583-A2.
30-JAN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                  Query Match
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AAK86282 standard; DNA; 65854 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:41094.
WO200157182-A2.
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PA (1515-) ISIS PHARM INC.

Query Match

Bast Local Similarity 90.0%; Pred. No. 1.9e-06;

ID ACN44342 pt-2-2-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PD 12-SEP-2003.
PA (SAGRE DISCOVERY.
Query Match 2.7%; Score 67.2; DB 11; Length 61103;
Best Local Similarity 90.0%; Pred. No. 1.7e-06;
RESULT 955
                                                                                                                                                                                                                                                                                                                                                                                                          Length 59554;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADG88330 standard; DNA; 66804 BP.

3. Human transporter protein genomic DNA.

2. 7%; Score 67.2; DB 10; Length 66804;
Best Local Similarity 90.0%; Pred. No. 1.8e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 59554;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.7%; Score 67.2; DB 11; Length 60381; 90.0%; Pred. No. 1.7e-06;
                                                                                                        Length 59554;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 73544;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 66804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 65854
                                                                                                                                                                                                                                                                                          PD 30-JAN-2003.
PA (SAGR-) SAGRES DISCOVERY.
Query Match
Best Local Similarity 90.0%; Pred. No. 1.7e-06;
RESULT 952
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADB9594 standard; DNA; 59554 BP.

Human TK2 gene genomic DNA sequence.

MO20030199484-A2.

15-MAY-2003.

(SAGR-) SAGRES DISCOVERY.

2.7%; Score 67.2; DB 10;

Bt Local Similarity 90.0%; Pred. No. 1.7e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 67.2; DB 4;
ery Match 90.0%; Pred. No. 1.7e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABK87050 standard; cDNA; 66804 BP.
Human transporter protein genomic DNA.
Query Match
2.7%; Score 67.2; DB 6;
Best Local Similarity 90.0%; Pred. No. 1.8e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.7%; Score 67.2; DB 8;
90.0%; Pred. No. 1.9e-06;
                                                                                                            Score 67.2; DB 9;
Pred. No. 1.7e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 958
ID ABS56296 standard; CDNA; 73544 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACN43886 standard; DNA; 61103 BP.
Human genomic sequence hCG1784975
WO2003073826-A2.
                                                                                                                                                                                                                    ADB72434 standard; DNA; 59554 BP.
Human TK2 gene.
WO2003008583-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ACN44494 standard; DNA; 60381 BP.
Human genomic sequence hCG27935.
WO2003073826-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADP66763 standard; DNA; 75033 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human transporter protein gene. WO200279252-A1.
                                                                                                            2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Deet in accordant of the following sequence hCG2 by MCO03073826-A2.

PA (SAREA) SAGRES DISCOVERY.

Query Match SAGRES DISCOVERY.

Query Match SAGRES DISCOVERY.

Deet Local Similarity 90.0%;

RESULT 954

ID ACM4388 standard; DNA, 611

DE Human genomic sequence hCG1

DE HUMAN GENOMIC SEQUENCE SEQUE
                                  17-JUL-2003.
(SAGR-) SAGRES DISCOVERY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-OCT-2002.
(PEKE ) PE CORP NY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                    Query Match
Best Local Similarity
RESULT 951
WO2003057146-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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Glutamine:fructose-6-phosphate amide transferase 1 genomic sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 86764;
                                                                                 Score 67.2; DB 11; Length 78268;
Pred. No. 1.9e-06;
                                                                                                                                                                                                                                                                                                                                                                                                  Length 79597;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.7%; Score 67.2; DB 10; Length 79652; 90.0%; Pred. No. 1.9e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.7%; Score 67.2; DB 12; Length 80928; 90.0%; Pred. No. 2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 88939;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 89182;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 93390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.7%; Score 67.2; DB 6; Length 98865; 90.0%; Pred. No. 2.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA51405 standard; DNA, 80959 BP.
Human secreted protein gene sequence, SEQ ID No 3.

Nuery Match
2.7%; Score 67.2; DB 8; Length 80959;
dest Local Similarity 90.0%; Pred. No. 2e-06;
                                                                                                                                                                                                                                           Length 78539;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABQ78054 standard; cDNA; 98865 BP.
Human Ras-like protein encoding genomic DNA SEQ ID NO
WO200261088-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AD097623 standard; DNA; 86764 BP.
Human cancer associated sequence HD10-016, SEQ ID 600.
WO2004060304-A2.
                                                                                                                                                                                                                                                                                                     ADL13866 standard; DNA; 79597 BP.
Osteoarthritis-associated polymorphic nucleotide #398.
WO2003054166-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADO25290 standard; DNA; 80928 BP.
Human protein kinase C iota/lambda genomic sequence.
WO2004041212-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 2.7%; Score 67.2; DB 10; Best Local Similarity 90.0%; Pred. No. 2.1e-06; RESULT 970
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 67.2; DB 12;
Pred. No. 2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 67.2; DB 13;
Pred. No. 2.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.7%; Score 67.2; DB 11; 90.0%; Pred. No. 2.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                       Score 67.2; DB 10;
Pred. No. 1.9e-06;
                                                                                                                                                                                                                                Query Match 2.7%; Score 67.2; DB 8;
Best Local Similarity 90.0%; Pred. No. 1.9e-06;
RESULT 962
                                                                                                                                       ACA64942 standard; DNA; 78539 BP.
Human FRAP1 DNA corresponding to AL049659
B10127372-A1.
05-DEC-2002.
(PATH-) PATHOARRAY GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA; 93390 BP
Human genomic sequence hCG17932.
WO2003073826-A2.
12-SEP-2003.
(SARES DISCOVERY.
Query Match
Best Local Similarity 90.0%; Pre
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(SAGRE) SAGRES DISCOVERY INC.

(SAGR.) SAGRES DISCOVERY INC.

2.7%;

lery Match 2.7%;
                                                                                                                                                                                                                                                                                                                                                     WOADCOCCE

GOUCK-) INCYTE GENOMICS INC.

Lery Match 2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAY-2004.
(SACK/) SACKTOR T C.
(CRAR/) CRARY J F.
(HERN/) HERNANDEZ A I.
(MIRR/) MIRRA S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-MAR-2003.
(SANY ) SANKYO CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
RESULT 966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
RESULT 967
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
RESULT 968
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Best Local Similarity
RESULT 969
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
RESULT 971
                                                                                                                                                                                                                                                                                                                                                                                                       Ouery Match
Best Local Similarity
RESULT 963
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-AUG-2002.
(PEKE ) PE CORP NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADD71350 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 965
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Best Local Similarity
RESULT 992
                 Best Local Similarity RESULT 982
                                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 985
                                                       Query Match
Best Local Similarity
RESULT 983
                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
RESULT 987
                                                                                                                                                                                                                                                                                                                                                                                                                        ADNO6353 standard;
Human FLAP genomic
WO2004035741-A2.
                                                                                                                             Best Local Similarity
RESULT 984
                                                                                                                                                                                                                                                                                                                              Best Local Similarity
RESULT 986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
RESULT 989
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Best Local Similarity
RESULT 993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
RESULT 990
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RESULT 991
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Best Local Similarity
RESULT 988
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AMHP ) WYETH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AD032260 standard; DNA; 106707 BP.
Human chromosome 20 region containing type 2 diabetes mellitus genes
                                                                                                                                                                                                                                                                                                                                                                                                                                        PD 15-JUL-2004.

PA (SAGR-) SAGRES DISCOVERY INC.

Query Match 2.7%; Score 67.2; DB 13; Length 101193;

Beet Local Similarity 90.0%; Pred. No. 2.2e-06;

RESULT 975
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.7%; Score 67.2; DB 13; Length 107036; 90.0%; Pred. No. 2.3e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.7%; Score 67.2; DB 12; Length 101270; 90.0%; Pred. No. 2.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .Query Match 2.7%; Score 67.2; DB 13; Length 106544; Best Local Similarity 90.0%; Pred. No. 2.3e-06; RESULT 977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.7%; Score 67.2; DB 12; Length 106707; 90.0%; Pred. No. 2.3e-06;
                                                                                             Length 99046;
                                                                                                                                   ID ADOS6274 standard; DNA; 99100 BP.
DB Human cyclin-dependent kinase 10, CDK10, genomic sequence.
Query Match 2.7%; .Score 67.2; DB 12; Length 99100;
Best Local Similarity 90.0%; Pred. No. 2.2e-06;
RESULT 973
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 107818,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 110000;
                                                                                                                                                                                                                                                                                                                                                  Length 99500;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADQ17814 standard; DNA; 101270 BP.
Human soft tissue sarcoma-upregulated DNA - SEQ ID
WO2004048938-A2.
                                                     15-JUL-2004.

1 (SAGR-) SAGRES DISCOVERY INC.

Query Match
2.7%; Score 67.2; DB 13;
Best Local Similarity 90.0%; Pred. No. 2.2e-06;
                                                                                                                                                                                                                                                                                                      04-JUN-2002.
(ISIS-) ISIS PHARM INC.
2.7%; Score 67.2; DB 6;
st Local Similarity 90.0%; Pred. No. 2.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 67.2; DB 4;
Pred. No. 2.3e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 67.2; DB 9;
Pred. No. 2.3e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABD32669 standard; DNA; 106544 BP.
Human cancer-associated genomic DNA HD13-073.
WO20040474320-A2.
(SAGR-) SAGRES DISCOVERY INC.
ABD33291 standard; DNA; 99046 BP.
Human cancer-associated (CA) gene HD07-051.
WO2004058146-A2.
                                                                                                                                                                                                                                                                                                                                                                                      Human cancer-associated (CA) gene HD07-068.
WQ2004058146-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABD33557 standard; DNA; 107036 BP.
Human cancer-associated (CA) gene HD07-111.
WO2004058146-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AALS7572 standard; cDNA; 107818 BP.
Human phosphomannomutase 1 cDNA.
WO2003059148-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (JOSL-) JOSLIN DIABETES CENT INC.
                                                                                                                                                                                                                                                DNA; 99500 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-JUN-2004.
(PROT-) PROTEIN DESIGN LABS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-JUL-2004.
(SAGR-) SAGRES DISCOVERY INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-JUL-2003.
(GENE-) GENE LOGIC INC.
(LGBI-) LG BIOMEDICAL INST.
                                                                                                                                                                                                                                                AAD41740 standard; DI
Human RECQL2 DNA #1.
US6399378-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
RESULT 980
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Best Local Similarity
RESULT 981
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related secreted protein.
                                                                                                                                                                                        Human transporter protein encoding gene SEQ ID NO:1.
2.7%; Score 67.2; DB 10; Length 110000; st Local Similarity 90.0%; Pred. No. 2.3e-06;
                                                                                                                                                                                                                                                                                                                                                             Length 110000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 110000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 110000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 110000;
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(GENE-) GENE LOGIC INC.
ery Match 2.7%; Score 67.2; DB 6; Length 110608;
                                                         Length 110000;
 Length 110000;
                                                                                                                 Length 110000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABK83572 standard; cDNA; 110608 BP.
Human cDNA differentially expressed in granulocytic cells #143.
WO200228999-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADQ97050 standard; DNA; 687411 BP.
Human cancer associated sequence HD08-001, SEQ ID 26.
WO2004060304-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human 5-lipoxygenase activating protein (FLAP) gene-WO2004035746-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.7%; Score 67.2; DB 12; 90.0%; Pred. No. 2.3e-06;
                                                                                                                                                                                                                                                               ALIS2246 standard; DNA; 378361 BP.
Human genomic DNA for the gene encoding the EGF r
Query March
2.74; Score 67.2; DB 10;
Best Local Similarity 90.0%; Pred. No. 2.3e-06;
                                                                                                                                                                                                                                                                                                                                                               2.7%; Score 67.2; DB 12; 90.0%; Pred. No. 2.3e-06;
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29-APR-2004.
(DECO-) DECODE GENETICS EHF.
(ery Match 2.7%; Score 67.2; DB 13;
ery Match ".""larity 90.0%; Pred. No. 2.3e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 67.2; DB 12;
Pred. No. 2.3e-06;
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Pred. No. 2.3e-06;
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Pred. No. 2.3e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 67.2; DB 12;
Pred. No. 2.3e-06;
 Score 67.2; DB 4;
Pred. No. 2.3e-06;
                                                       Score 67.2; DB 6;
Pred. No. 2.3e-06;
                                                                                                                 Score 67.2; DB 6;
Pred. No. 2.3e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Drug therapy altered expressed gene #193
WO2004072265-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                        DNA; 398800 BP.
DNA SEQ ID NO:1
                                                                                                                                                                      ADE11169 standard; DNA; 394191 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADS94372 standard; DNA; 398800 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA; 110608
                                                                                                               2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OBCO-) DECODE GENETICS EHF.

(BECO-) DECODE GENETICS EHF.

(ery Match ....larity 90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.78;
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2.7%;
                                                       2.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.7%;
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Length 117829;

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Human Na+independent transporter-related transporter protein gene. rry Match 2.7%; Score 67.2; DB 10; Length 139573; it Local Similarity 90.0%; Pred. No. 2.6e-06;
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RESULT 1008
ID ADJ96282 standard; DNA; 130001 BP.
DE Human breast cancer-1 DNA #14.
PN US2004014051-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    vuery Match 2.7%; Score 67.2; DB 12; Length 130001;
Best Local Similarity 90.0%; Pred. No. 2.5e-06;
RESULT 1009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 67.2; DB 11; Length 133632;
Pred. No. 2.6e-06;
                                                                                                                                               2.7%; Score 67.2; DB 12; Length 117143; 90.0%; Pred. No. 2.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 2.7%; Score 67.2; DB 12; Length 120447; Best Local Similarity 90.0%; Pred. No. 2.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-SEP-2003.
(SAGR.) SAGRES DISCOVERY.
2.7%; Score 67.2; DB 11; Length 122923;
Rt Local Similarity 90.0%; Pred. No. 2.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 2.7%; Score 67.2; DB 6; Length 130263;
Best Local Similarity 90.0%; Pred. No. 2.5e-06;
RESULT 1010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABK83573 standard; cDNA; 130263 BP.
Human cDNA differentially expressed in granulocytic cells #144.
WO200228999-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABQ88154 standard, cDNA, 147708 BP.
Human osteoblast differentiation related cDNA SEQ ID NO 61.
87-070250301-A2.
                                                                                                                                                                                                                                                                                                                                                              ADQ97391 standard; DNA; 120447 BP.
Human cancer associated sequence HD08-042, SEQ ID 368.
MO2004060304-A2.
22-JUL_2004.
(SAGR-) SAGRES DISCOVERY INC.
                                                                                                                                                                                                ADQ97319 standard; DNA; 117829 BP.
Human cancer associated sequence HD08-031, SEQ ID 296.
WO2004060304-A2.
                                                                                                                                                                                                                                                      PD 22-JUL-2004.
PA (SAGR-) SAGRES DISCOVERY INC.
Query Match
Best Local Similarity 90.0%; Pred. No. 2.4e-06;
RESULT 1005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADRS2731 standard; DNA; 127917 BP.
Drug therapy altered expressed gene #82
WO2004072265-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADH58564 standard; DNA; 139573 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACN45054 standard; DNA; 133632 BP.
Human genomic sequence hCG28560.
WO2003073826-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .T 1006
ACN44026 standard; DNA; 122923 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human genomic sequence hCG24510.
WO2003073826-A2.
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12-SEP-2003.
(SAGR-) SAGRES DISCOVERY.
PART ATCh 2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (AMHP) WYETH.
(BURC/) BURCZYNSKI M.
(TWIN/) TWINE N.
(DORN/) DORNER A J.
(TREP/) TREPICCHIO W L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-APR-2002.
(GENE-) GENE LOGIC INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      22-JAN-2004.
(ISIS-) ISIS PHARM INC.
                                                                                                                               (GENO-) GENOX RES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
RESULT 1011
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Best Local Similarity
RESULT 1012
                                                                                                                                                                Best Local Similarity RESULT 1004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                   Query Match
Best Local Similarity 90.0%; Score 67.2; DB 4; Length 114793; RESULT 1001

1D ACD13448 standard; DNA; 115756 BP.
DB Human DNA encoding a p53 modifi**
PD 12-DEC-200*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 67.2; DB 13; Length 115935;
Pred. No. 2.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.7%; Score 67.2; DB 13; Length 113604; 90.0%; Pred. No. 2.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 2.7%; Score 67.2; DB 13; Length 113604; Best Local Similarity 90.0%; Pred. No. 2.4e-06;
                                                                                                                                                                                                                                                                                                                                  Length 113604;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 115756;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 113604;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADS19854 standard; DNA; 113604 BP.
Human chromosome 21 nucleotide sequence SEQ ID NO:2.
WO2004018701-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADS19853 standard; DNA; 113604 BP.
Human chromosome 21 nucleotide sequence SEQ ID NO:1.
WO2004018701-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               vuery Match
Beet Local Similarity 90.0%; Pred. No. 2.4e-06;
RESULT 1002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Aatch 2.7%; Score 67.2; DB 9; Local Similarity 90.0%; Pred. No. 2.4e-06;
                                                                                                                                                                                                                                                                                                                            Query Match 2.7%; Score 67.2; DB 9; Best Local Similarity 90.0%; Pred. No. 2.4e-06; RESULT 997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADR52890 standard; DNA; 115935 BP.
Drug therapy altered expressed gene #241.
WO2004072265-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADJ74882 standard; DNA; 117143 BP.
Marker gene SEQ ID NO:134.
EP1194274-A2.
03-MAR-2004.
                                                                                                                                                                                                                                                                                                                                                                                     ADA42606 standard; DNA; 113604 BP.
Human LQTS genomic DNA #1.
US2003077633-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90.06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (PERL-) PERLEGEN SCI INC. (COXD/) COX D R. (ARNO/) ARNOLD D A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-APR-2003.
(PERL-) PERLEGEN SCI INC.
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BURCZYNSKI M.
(TWIN/) TWINE N.
(DORN/) DORNER A J.
(TREP/) TREPICCHIO W L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ocal Similarity
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Best Local Si
RESULT 999
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Length 161425;

Length 162025;

Length 162025;

Length 162025;

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ADS75958 standard; DNA; 162025 BP.
Human A-Kinase anchoring protein 10 allele AKAP10-6 genomic sequence.
W02004081576-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADS75959 standard; DNA; 162025 BP.
Human A-kinase anchoring protein 10 allele AKAP10-7 genomic sequence.
WO2004081576-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human A-kinase anchoring protein 10 allele AKAP10-1 genomic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.7%; Score 67.2; DB 13; Length 162025; 90.0%; Pred. No. 2.9e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 67.2; DB 10; Length 170834; Pred. No. 3e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.7%; Score 67.2; DB 13; Length 162025; 90.0%; Pred. No. 2.9e-06;
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Human A-kinase anchoring protein 10 wild type genomic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAD62833 standard; DNA; 170834 BP.
Human BAC #2 containing formin (Fmn)-2 genomic DNA
US2003170683-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Jr 1030
Bable8122 standard; DNA; 174424 BP.
Ovary cancer related gene sequence SEQ ID NO:6459.
WO200194629-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 67.2; DB 13;
Pred. No. 2.9e-06;
                                            2.7%; Score 67.2; DB 4; 90.0%; Pred. No. 2.9e-06;
                                                                                                                                                                                                                     2.7%; Score 67.2; DB 4;
90.0%; Pred. No. 2.9e-06;
                                                                                                                                                                                                                                                                                                                                                                                              Score 67.2; DB 6;
Pred. No. 2.9e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 67.2; DB 6;
Pred. No. 2.9e-06;
                                                                                                                                                                                                                                                                                                      Human AKAP allelic variant (AKAP10-6) gene.
WO200204489-A2.
17-JAN-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAD28759 standard; DNA; 162025 BP.
Human AKAP allelic variant (AKAP10-7) gene.
WO200204489-A2.
                                                                                                     AAH02339 standard; DNA; 162025 BP.
Human AKAP10 gene SEQ ID NO: 35.
MO200127857-A2.
19-APR-2001.
(SEQU-) SEQUENOM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADS75963 standard; DNA; 162025 BP
                                                                                                                                                                                                                                                                                      DNA; 162025
                                                                                                                                                                                                                                                                                                                                                                                              2.7%;
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                        SEQUENOM INC.
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(SEQU-) SEQUENOM INC.
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(SEQU-) SEQUENOM INC.
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(SEQU-) SEQUENOM INC.
                                            Query Match
Best Local Similarity
RESULT 1022
                                                                                                                                                                                                                                         Local Similarity
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Best Local Similarity
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Best Local Similarity
RESULT 1029
                                                                                                                                                                                                                                                                                      AAD28758 standard;
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      19-APR-2001
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PA (PROT-) PROTEIN DESIGN LABS INC.

QUERTY MATCH

2.7%; Score 67.2; DB 12; Length 155937;

Best Local Similarity 90.0%; Pred. No. 2.8e-06;

RESULT 1018
                                                                                                                                                                                                                   Length 152759;
                                                                                                                                                                                                                                                                                                                                                                                           Length 153752;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 154465;
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                                          Length 147708;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADQ19389 standard; DNA; 155937 BP.
Human soft tissue sarcoma-upregulated DNA - SEQ ID 2208
WO2004048938-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABQ88179 standard; cDNA; 160771 BP.
Human osteoblast differentiation related cDNA SEQ 1D NO
WQ200250301-A2.
                                                                                                                                                                                                                                                                           ADQ97531 standard; DNA; 153752 BP.
Human cancer associated sequence HD09-009, SEQ ID 508.
WO2004060304-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADL13861 standard; DNA; 154681 BP.
Osteoarthritis-associated polymorphic nucleotide #393.
                                                                                                                                                                                                                       DB 13;
                                                                                                                                                                                                                                                                                                                                                                                           2.7%; Score 67.2; DB 12;
90.0%; Pred. No. 2.8e-06;
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(INCY-) INCYTE GENOMICS INC.
2.7%; Score 67.2; DB 10;
ery Match....iarity 90.0%; Pred. No. 2.8e-06;
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                             Query Match 2.7%; Score 67.2; DB 6; Best Local Similarity 90.0%; Pred. No. 2.7e-06; REGULT 1013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 2.7%; Score 67.2; DB 6; Best Local Similarity 90.0%; Pred. No. 2.8e-06; RESULT 1016
                                                                                                     ID ABD33075 standard; DNA; 152759 BP.

DE Human cancer-associated (CA) gene HD07-002.

PN WC2004058146-A2.

PD 15-JUL-2004.

PA (SAGR-) SAGRES DISCOVERY INC.

QUESTY MARCH

Best Local Similarity 90.0%; Pred. No. 2.8e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABD32817 standard; DNA; 156416 BP.
Human cancer-associated genomic DNA HD17-001.
02-SEP-2004.
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Human gene associated with low HDL-C FABP-3.
US2004043389-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAD28763 standard; DNA; 154465 BP.
Human AKAP allelic variant (AKAP10-1) gene.
WO200204489-A2.
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Human AKAP10 gene SEQ ID NO: 36.
WO200127857-A2.
                                                                                                                                                                                                                                                                                                                                              22-JUL-2004.
(SAGR-) SAGRES DISCOVERY INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SAGR-) SAGRES DISCOVERY INC.
(GENE-) GENE LOGIC INC.
(PROC ) PROCTER & GAMBLE CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GENE-) GENE LOGIC INC.
(PROC ) PROCTER & GAMBLE CO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-JAN-2002.
(SEQU-) SEQUENOM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-MAR-2004.
(VITI-) VITIVITY INC.
                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity RESULT 1015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 1019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity RESULT 1017
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Length 162025

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2.7%; Score 67.2; DB 11; Length 198522; 90.0%; Pred. No. 3.2e-06;
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(SAGR-) SAGRES DISCOVERY INC.
                                                                            (SAGR-) SAGRES DISCOVERY.
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(VITI-) VITIVITY INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
RESULT 1045
                                                                                                                                                                                                                                                                                                                                                                                                                                  Best_Local Similarity RESULT 1042
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                                                                                              Query Match
Best Local Similarity
RESULT 1040
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RESULT 1047
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                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                              RESULT 1041
Best Local Similarity 90.0%; Pred. No. 3e-06; Length 174424; RESULT 1031

ID ACN43946 standard; DNA; 174448 BP. BP HWO2003073826-A2.

PD 12-SEP-200?
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AD097687 standard; DNA; 179487 BP.

Human cancer associated sequence HD10-028, SEQ ID 664.

Human cancer associated sequence HD10-028, SEQ ID 664.

22-JUL-2004.

(SAGR.) SAGRES DISCOVERY INC.

2.7%; Score 67.2; DB 12; Length 179487; St Local Similarity 90.0%; Pred. No. 3e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         wwery match 2.7%; Score 67.2; DB 12; Length 181343; Best Local Similarity 90.0%; Pred. No. 3.1e-06; RESULT 1038
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-JUL-2004.
(GAGR-) SAGRES DISCOVERY INC.
ery Match
2.7%; Score 67.2; DB 12; Length 194534;
st Local Similarity 90.0%; Pred. No. 3.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-JUN-2004.

(PROT-) PROTEIN DESIGN LABS INC.

ery Match 2.7%; Score 67.2; DB 12; Length 175737;

-- Tocal Similarity 90.0%; Pred. No. 3e-06;
                                                                                                                                                                                                                                                                              11; Length 174448;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 175077;
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(GENE-) GENE LOGIC INC.
ery Match 2.7%; Score 67.2; DB 6; Length 175737;
ery Match 90.0%; Pred. No. 3e-06;
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ABR83571 standard; cDNA; 175737 BP.
Human CDNA differentially expressed in granulocytic cells #142
WO200228999-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADO18934 standard, DNA; 175737 BP.
Human soft tissue sarcoma-upregulated DNA - SEQ ID 1753
WO2004048938-A2.
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E Human soft tissue sarcoma-upregulated DNA - SEQ ID 2392.

N W02004048938-A2.

D 10-JUN-2004.

A (PROT-) PROTEIN DESIGN LABS INC.

Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ALL13596 standard; DNA; 175737 BP.
Osteoarthritis-associated polymorphic nucleotide #128.
WO2003054166-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADQ97481 standard; DNA; 194534 BP.
Human cancer associated sequence HD09-002, SEQ ID 458
WO2004060304-A2.
                                                                                                                                                                                                                                                                                                                                                                           PD 12-SEP-2003.
PA (SAGR-) SAGRES DISCOVERY.
Query Match
Best Local Similarity 90.0%; Pred. No. 3e-06;
RESULT 1033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            O3-JUL-2003.
(INCY-) INCYTE GENOMICS INC.
(Ery Match
ery Match
?: "milarity 90.0%; Pred. No. 3e-06;
                                                                                                                                                                                                                                                                            Ouery Match 2.7%; Score 67.2; DB Best Local Similarity 90.0%; Pred. No. 3e-06; RESULT 1032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACN44010 standard; DNA; 198522 BP. Whuman genomic sequence hCG1643869; WOZ003073826-A2.
                                                                                                                                                                                                                                                                                                                                   ACN44626 standard; DNA; 175077 BP.
Human genomic sequence hCG19724.
WO2003073826-A2.
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JA-A2.

JR-) SAGRES DISC

LY MATCH

BESULT 1037

ID AD019573 stand**

PN W02004**

PD 10-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ouery Match
Best Local Similarity
RESULT 1034
ID ADL13596 standard, DN
DE Osteoarthritis-associ
PN W02003054466-A2.
PD 03-JUL-2003.
PA (INCY-) INCYTE GENOMI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
RESULT 1035
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AAV57903 standard; DNA; 237326 BP.
Hereditary haemochromatosis subregion from an HH affected individual.
                                                                                                                                                                                   ______AAV57926 standard, DNA, 235033 BP.
AAV57926 standard, DNA, 235033 BP.
Hereditary haemochromatosis subregion from an unaffected individual WO9814466-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EBET Local Similarity 90.0%; Pred. No. 3.7e-06; RESULT 1048
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AD059422 standard; DNA; 246940 BP.

Human cancer-associated (CA) gene sequence SEQ ID NO:58.

N WO2004089288-Al.

O 15_JUL_2004.

A (SAGR-) SAGRES DISCOVERY INC.

Query Match

Best Local Similarity 90.0%; Pred. No. 3.6e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SACR-) SAGRES DISCOVERY.

ry Match
2.7%; Score 67.2; DB 11; Length 220224;
t Local Similarity 90.0%; Pred. No. 3.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 247682;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PD 27-FEB-2003.

PA (MCLA-) MCLAUGHLIN RES INST.
Query Match 2.7%; Score 67.2; DB 8; Length 249999;
Query Match 2.7%; Pred. No. 3.7e-06;
RESULT 1046
                                                                                                                                                                                                                                                                                                                        Length 235033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 237326,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABZ80229 standard; DNA; 249999 BP.
Human tramdorin gene region genomic DNA SEQ ID NO:26.
WO2003016502-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.7%; Score 67.2; DB 12; 90.0%; Pred. No. 3.6e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                              PD 09-APR-1998.
PA (PROG-) PROGENTIOR INC.
2.7%; Score 67.2; DB 2;
Best Local Similarity 90.0%; Pred. No. 3.6e-06;
                                                                                                                                                                                                                                                             WOJOGENIUS

PO-ACACH PROGENTIOR INC.

(PROGENIUS 178; Score 67.2; DB 2; LETY MATCH 2.7%; Score 67.2; DB 2; LETY MATCH 90.0%; Pred. No. 3.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABD32602 standard, DNA; 277616 BP.
Human cancer-associated genomic DNA HD7-221.
W02004074320-A2.
(SAGR-) SAGRES DISCOVERY INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABD33570 standard; DNA; 256157 BP.
Human cancer-associated (CA) gene HD07-114.
WO2004058146-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADLOBIO9 standard; DNA; 247682 BP.
Human gene associated with low HDL-C AT3
US2004043389-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACN44650 standard; DNA; 256157 BP.
Human genomic sequence hCG38672.
WO2003073826-A2.
ACN44702 standard; DNA, 220224 BP.
Human genomic sequence hCG1788599
WC2003073826-A2.
12-5EP-2003.
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27-DEC-2001
(GSFU-) GSF
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(DMAS/)
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:37166.
WO200157182-A2.
2.7%; Score 67.2; DB 13; Length 277616; 90.0%; Pred. No. 3.9e-06;
                                                                                                                  15-JUL-2004.
(SAGR-) SAGRES DISCOVERY INC.
(SAGR-) SAGRES DISCOVERY INC.
2.7%; Score 67.2; DB 12; Length 299598; Bt Local Similarity 90.0%; Pred. No. 4e-06;
                                                                                                                                                                                                                                                                                                    Query Match 2.7%; Score 67.2; DB 12; Length 312477; Best Local Similarity 90.0%; Pred. No. 4.1e-06; RESULT 1051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.
(ADSC.) CENT ADVANCED SCI & TECHNOLOGY INCUBATIO.

FY MAICH

L Local Similarity 90.0%; Pred. No. 4.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Match 2.7%; Score 67.2; DB 4; Length 325791; Local Similarity 90.0%; Pred. No. 4.2e-06;
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(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 67; DB 4; Length 4164;
st Local Similarity 88.0%; Pred. No. 4.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAKS8620 standard, DNA, 14646 BP.
Human HHV4 DNA replication inhibitor TB7 coding sequence #1
WO200198489-A1.
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                                       ADOS9380 standard; DNA; 299598 BP.
Human cancer-associated (CA) gene sequence SEQ ID NO:16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABS67550 standard; DNA; 7219 BP.
Novel human DNA repair protein, genomic sequence #33
US2002086353-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Aurijji standard; DNA; 1004 BP.
Human Coproporphyrinogen oxidase DNA, SEQ ID 26.
WO2003012813-A2.
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88.0%; Pred. No. 5.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                    Human Oestrogen receptor beta gene WO200162793-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADC8616 standard; DNA; 349989 BP.
Human GPCR gene SEQ ID NO:1369.
EP1270724-A2.
                                                                                                                                                                                                                                                                                                                                                                                 DNA; 325791 BP.
                                                                                                                                                                                                                       DNA; 312477 BP
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                                                                                                                                                                                                               ADP69744 standard; DNA; 3
Human ROCK 1 DNA #2.
US2004115641-A1.
17-JUN-2004.
(ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PEKE ) PE CORP NY.
Query Match
Best Local Similarity
RESULT 1049
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ROSE/) ROSEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                                                                                                                 AAS43104 standard;
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(FARB ) BAYER AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-JAN-2003
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RESULT 1
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Human sequence from clone RPS-1009E24 on chromosome 20 Contains DNA WO2003072827-A1.
                                                                                                                                                                                                                                                        ABL97583 standard; DNA; 32249 BP.
Human testicular antigen encoding DNA fragment SEQ ID NO: 2235.
WO200155317-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 161484;
                                                                                                       Human reproductive system related antigen DNA SEQ ID NO: 7364
WO200155320-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.7%; Score 67; DB 8; Length 207433; 73.9%; Pred. No. 3.7e-06;
PA (GSFU-) GSF FORSCHUNGSZENTRUM UMWELT & GESUNDHEI.

Query Match
2.7%; Score 67; DB 6; Length 14646;
Best Local Similarity 88.0%; Pred. No. 8.6e-07;
RESULT 1058
ID AAL04676 standard; DNA; 32249 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 57038
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                                                                                                                                                                                          Length 32249;
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Human sequence from clone RP5-1009E24 on chromosome 20
WO2003072827-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABZ72040 standard; DNA; 207433 BP.
Gene 216 H194BAC1098L22 nucleotide sequence SEQ ID NO W0200178894-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-SEP-2003.
(CHIL-) CHILDREN'S HOSPITAL MEDICAL CENT.
ery Match
2.7%; Score 67; DB 11;
                                                                                                                                                                  (HUMA-) HUMAN GENOME SCI INC.

ry Match 2.7%; Score 67; DB 4; 1

t Local Similarity 88.0%; Pred. No. 1.3e-06;
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73.9%; Pred. No. 3.7e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   04-SEP-2003.
(CHIL-) CHILDREN'S HOSPITAL MEDICAL CENT.
ery Match
2.7%; Score 67; DB 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.7%; Score 67; DB 13; 93.3%; Pred. No. 3.3e-06;
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88.0%; Pred. No. 1.3e-06;
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Bacterial artificial chromosome RPCI-11
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BAC1098L22 DNA sequence.
WO200283077-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-OCT-2002.
(SCHE ) SCHERING CORP.
(GENO-) GENOME THERAPEUTICS CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GENO-) GENOME THERAPEUTICS CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                            ADP65343 standard; DNA; 57038 BP
                                                                                                                                                                                                                                                                                                                                       HUMAN GENOME SCI INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-JUN-2004.
(ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DUPUIS J.
DEL MASTRO R
                                                                                                                                                                                                     Best Local Similarity
RESULT 1059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity RESULT 1061
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                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
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RESULT 1064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADS19665 standard;
Human PTPN13 cDNA.
US2004115635-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SIMON J.
ALLEN K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JS2004002470-A1.
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(ALLE/)
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Human gene associated with low HDL-C LRPAPI.
US2004043389-A1.
                                    04-MAR-2004.
(VITI-) VITIVITY INC.
                                                                                 Query Match
Best Local Similarity
RESULT 1075
                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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1077
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18-AUG-2004,
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:25082.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAK78813 standard; DNA; 51469 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:33625.
WO200157182-A2.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:24134
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 34722;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             vuery Match 2.7%; Score 66.8; DB 9; Length 98829;
Best Local Similarity 86.0%; Pred. No. 2.8e-06;
RESULT 1074
                                                                                                                                                                                                                                                                                                                                      2.7%; Score 67; DB 12; Length 207433; 73.9%; Pred. No. 3.7e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-JUN-2004.
(PROT-) PROTEIN DESIGN LABS INC.
ery Match 2.7%; Score 66.8; DB 12; Length 3820;
st Local Similarity 86.0%; Pred. No. 4.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ID ABQ82235 standard; DNA; 43543 BP.

DE Human phospholipase protein encoding DNA SEQ ID NO:3.

Query Match 2.7%; Score 66.8; DB 6; Length 43543;

Best Local Similarity 78.3%; Pred. No. 1.8e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 66.8; DB 4; Length 51469;
st Local Similarity 86.0%; Pred. No. 2e-06;
                       2.7%; Score 67; DB 12; Length 207433; 73.9%; Pred. No. 3.7e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 66.8; DB 4; Length 51469;
8t Local Similarity 86.0%; Pred. No. 2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               O 99-AUG-2001.

A (HUMA-) HUMAN GENOME SCI INC.
Query Match 2.7%; Score 66.8; DB 4; Length 51469;
Best Local Similarity 86.0%; Pred. No. 2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                     Human soft tissue sarcoma-upregulated DNA - SEQ ID 5474
WO2004048938-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-JUL-2004.

4 (SAGR-) SAGRES DISCOVERY INC.

2.7%; Score 66.8; DB 13

Pest Local Similarity 91.0%; Pred. No. 1.6e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABD33520 standard; DNA; 34722 BP.
Human cancer-associated (CA) gene HD07-102.
WO2004058146-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAD55116 standard; DNA; 98829 BP.
Human ACE reference DNA (GI 13027555).
WO2003020118-A2.
                                                                                     ADL81193 standard; DNA; 207433 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADL08124 standard; DNA; 176080 BP
                                                                                                                                                                                                                                                                                                                                                                                                  ADQ22654 standard; DNA; 3820 BP.
                                                                                                                                                                                                                          (DUAS/) DUEL MASTRO R G. (SINO/) SINON J. (ALLE) ALLEN K. (PAND/) PANDIT S.
                                                                                                           BAC1098L22 DNA sequence
US2004023215-A1.
                                                                                                                                                                   KEITH T.
LITTLE R D.
EERDEWEGH P V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13-MAR-2003.
(VITI-) VITIVITY INC.
                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
RESULT 1067
                                      Best Local Similarity RESULT 1066
      (PAND/) PANDIT S.
  PA (PANE, , Ouery Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Si
RESULT 1068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Si
RESULT 1071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                        (LITT/)
(EERD/)
(DUPU/)
(DMAS/)
(SIMO/)
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ADRO8411 standard; cDNA; 3923 BP.
Full length human cDNA useful for treating neurological disease Seq 1917.
EP1447413-A2.
Length 176080;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 2235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.7%; Score 66.6; DB 3; Length 792; 88.9%; Pred. No. 2.2e-07;
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                                                                AAC72115 standard; DNA; 792 BP.
Single nucleotide polymorphism containing sequence #648
WOZ00058519-A2.
05-OCT-2000.
                                                                                                                                                                                                                                                               AAC72109 standard; DNA; 792 BP.
Single nucleotide polymorphism containing sequence #646
WO200058519-A2.
05-OCT-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAC72079 standard; DNA; 792 BP.
Single nucleotide polymorphism containing sequence #636.
WO200058519-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAC72094 standard; DNA; 792 BP.
Single nucleotide polymorphism containing sequence #641
WO200058519-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAC72112 standard; DNA; 792 BP.
Single nucleotide polymorphism containing sequence #647.
WO200058519-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAC72106 standard; DNA; 792 BP.
Single nucleotide polymorphism containing sequence #645.
WO200058519-A2.
  Score 66.8; DB 12;
Pred. No. 3.9e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.7%; Score 66.6; DB 3;
88.9%; Pred. No. 2.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                               2.7%; Score 66.6; DB 3;
38.9%; Pred. No. 2.2e-07;
                                                                                                                                                                                                 Score 66.6; DB 3;
Pred. No. 2.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.7%; Score 66.6; DB 3;
88.9%; Pred. No. 2.2e-07;
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Pred. No. 3.9e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 66.6; DB 3;
Pred. No. 2.2e-07;
                                                                                                                                                       (WHED ) WHITEHEAD INST BIOMEDICAL RES. (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (WHED ) WHITEHEAD INST BIOMEDICAL RES. (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (WHED ) WHITEHEAD INST BIOMEDICAL RES. (AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-OCT-2000.
(WHED ) WHITEHEAD INST BIOMEDICAL RES.
(AFFY-) AFFYMETRIX INC.
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(WHED ) WHITEHEAD INST BIOMEDICAL RES.
(AFFY-) AFFYMETRIX INC.
                                                                                                                                                                                                                                                                                                                                                        (WHED ) WHITEHEAD INST BIOMEDICAL RES. (AFFY-) AFFYMETRIX INC.
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Human polynucleotide SEQ ID NO 12840.
WO200164835-A2.
  2.7%;
86.0%;
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Length 7260;

Length 7476;

61

us-10-002-967a-58.rng.spdi

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AAK78286 standard; DNA; 28420 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:33098.
WO200157182-A2.
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PA (SAGR-) SAGRES DISCOVERY.
Query Match 2.7%; Score 66.6; DB 11; Length 22021;
Best Local Similarity 88.9%; Pred. No. 1.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PD 12-SEP-2003.
PA (SAGR.) SAGRES DISCOVERY.
Query Match 2.7%; Score 66.6; DB 11; Length 31116;
Best Local Similarity 88.9%; Pred. No. 1.7e-06;
RESULT 1097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Office States of The States of State
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Genomic sequence #416 encoding novel human enzyme polypeptide WO200155301-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 28420,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABK11076 standard; DNA; 35973 BP.
Human amyloid beta precursor protein binding protein 1 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAS96607 standard; DNA; 51552 BP.
DNA encoding human telomerase reverse transcriptase (TERT)
WO200188198-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human TCOF1 carcinoma associated gene, SEQ ID NO:1058
WO2003057146-A2.
                          Query Match 2.7%; Score 66.6; DB 11;
Best Local Similarity 88.9%; Pred. No. 7.5e-07;
                                                                                                                                                     ABB32850 standard; cDNA; 7476 BP.
Human cancer-associated cDNA HR17-016.2.
W02004074320-A2.
O2-SEP-2004.
COSAGRES DISCOVERY INC.
2.7%; Score 66.6; DB 13;
Best Local Similarity 88.9%; Pred. No. 7.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 66.6; DB 4;
ery Match 2.7%; Pred. No. 1.6e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.7%; Score 66.6; DB 6;
88.9%; Pred. No. 1.8e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-NOV-20U1.
(ISIS-) ISIS PHARM INC.
1ry Match 2.7%; Score 66.6; DB 6;
rry Match 88.9%; Pred. No. 2.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 66.6; DB 9;
Pred. No. 2.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACN44670 standard; DNA; 22021 BP.
Human genomic sequence hCG20786.
WO2003073826-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                 AAS42100 standard; DNA; 12595 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACN44954 standard; DNA; 31116 BP.
Human genomic sequence hCG38622.
WO2003073826-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADB72278 standard; DNA; 58822 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SAGR-) SAGRES DISCOVERY.

ry Match
t Local Similarity 88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
RESULT 1094
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SULT 1096
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABK13076 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
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WO2003008583-A2.
30-JAN-2003.
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10-JAN-2002.
      04-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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Human insulin-like growth factor 1 (somatomedin C) (IGF1), DNA sequence.
WO2003072827-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADG33110 standard; DNA; 7260 BP.
Human DNA differentially expressed in patients with SLE SeqID434.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK84583 standard; cDNA; 7260 BP.

Human cDNA differentially expressed in granulocytic cells #1154
W020022899-A2.
11.-APR-2002.
(GENE JOGIC INC.
2.7%; Score 66.6; DB 6; Length 7260;
ELocal Similarity 88.9%; Pred. No. 7.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABT11091 standard; cDNA; 7260 BP.
Human breast cancer associated coding sequence SEQ ID NO: 1225.
WO200259271-A2.
                             Match 2.7%; Score 66.6; DB 13; Length 3923;
Local Similarity 88.9%; Pred. No. 5.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 7260
                                                                                                                                                                                                                                                                                   Length 7260;
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                                                                                                                                                                                                                                                                                                                                                            PD 06-NOV-2003.

PA (EXPR.) EXPRESSION DIAGNOSTICS INC.

Query Match

2.7%; Score 66.6; DB 10;

Best Local Similarity 88.9%; Pred. No. 7.5e-07;

RESULT 1091
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WOLVELLOOD.

(GENE) GENE LOGIC INC.

(ery Match

2.7%; Score 66.6; DB 6;

(ery Match

2.7%; Pred. No. 7.5e-07;
                                                                                                   DE. Human endometrial cancer related gene, IGF1.

DE. Human endometrial cancer related gene, IGF1.

PN W0200209573-A2.

PN GGHM ) BRIGHAM & WOMENS HOSPITAL INC.

Query Match

Best Local Similarity 88.9%; Pred. No. 7.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 66.6; DB 6;
Pred. No. 7.5e-07;
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Query Match
2.7%; Score 66.6; DB 6;

Best Local Similarity 88.9%; Pred. No. 7.5e-07;

RESULT 1089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 66.6; DB 6;
Pred. No. 7.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.7%; Score 66.6; DB 9;
88.9%; Pred. No. 7.5e-07;
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Human insulin-like growth factor-I (IGF-I) DNA
US2003072761-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABK64812 standard; DNA; 7260 BP.
Human benign prostatic hyperplasia gene #707.
WO200212440-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABN97244 standard; DNA; 7260 BP.
Gene #3742 used to diagnose liver cancer.
WO200229103-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (BGHM ) BRIGHAM & WOMENS HOSPITAL INC.
(BAAK/) BAAK J.
(REAS-) RES ASSOC BIOTECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Match 2.7%;
Local Similarity 88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GENE-) GENE LOGIC INC.
(NISB ) JAPAN TOBACCO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity RESULT 1085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-APR-2003.
(LEBO/) LEBOWITZ J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
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Length 35973;

#1.

Length 51552;

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Genomic sequence #174 encoding novel human connective tissue polypeptide. WO200155343-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK42274 standard; DNA; 6265 BP.
Genomic sequence #173 encoding novel human connective tissue polypeptide.
W0200155343-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1117
ID AAK69531 standard, DNA, 10989 BP.
DE Human immune/haematopoietic antigen genomic sequence SEQ ID NO:24343.
PN WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human immune/haematopoietic antigen genomic sequence SEQ ID NO:19750.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 10989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.7%; Score 66.4; DB 9; Length 6265; 71.0%; Pred. No. 7.9e-07;
                                                                                                                                                                                                                                                                                                              PA (HELI-) HELIX RES INST.
2.7%; Score 66.4; DB 4; Length 3441;
Best Local Similarity 71.0%; Pred. No. 5.6e-07;
RESULT 1111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OZ-AUG-2001.
(HUWA-) HUMAN GENOME SCI INC.
(ery Match 1994; Length 6265;
(ery Match imilarity 71.0%; Pred. No. 7.9e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 8942;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 5629;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 9883
                                                                                                                         Length 491;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABQ73765 standard; cDNA; 9883 BP.
Human colon specific nucleic acid (CSNA) SEQ ID NO:71.
WO200248370-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USACATA 2003.
20-MAR-2003.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 66.4; DB 9;
lery Match 2.7%; Pred. No. 7.4e-07;
                                                                                                                         2.7%; Score 66.4; DB 9;
71.0%; Pred. No. 1.9e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-Aug-2001.
(HUDA-) HUMAN GENOME SCI INC.
(HUDA-) HUMAN GENOME SCI INC.
2.7%; Score 66.4; DB 4;
ery Match 2.7%; Pred. No. 7.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WOZOWIA

09-AUG-2001.

(HUMAL) HUMAN GENOME SCI INC.

2.7%; Score 66.4; DB 4;

lery Match

2.7%; Score 66.4; DB 4;

2.7%; Pred. No. 1.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 2.7%; Score 66.4; DB 6;
Best Local Similarity 74.2%; Pred. No. 1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.7%; Score 66.4; DB 5;
87.5%; Pred. No. 9.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADBG0430 standard, DNA; 6265 BP.
Connective tissue related genomic DNA #173.
US200304375-A1.
20-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF22938 standard; DNA; 8942 BP.
Atopy related gene sequence SEQ ID 2.
WO200104302-A1.
                                                                                                                                                                                                      AAH18185 standard; cDNA, 3441 BP.
Human cDNA sequence SEQ ID NO:18094.
EP1074617-A2.
07-FEB-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAK64938 standard; DNA; 19183 BP
                                                                                                                                                                                                                                                                                                                                                                                                                          ABK42275 standard; DNA; 5629 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HUMA-) HUMAN GENOME SCI INC.
Query Match 2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MITU ) MITSUBISHI CHEM CORP.
              (LABA/) LABAT I.
(STAC/) STACHE-CRAIN B.
(DICK/) DICKSON M C.
(JONE/) JONES L W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (DIAD-) DIADEXUS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ouery Match
Best Local Similarity
RESULT 1112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity RESULT 1114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 1115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                             Best Local Similarity RESULT 1110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-JAN-2001
                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAK87645 standard; DNA; 86 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:42457.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 104900;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 156416;
(SAGR-) SAGRES DISCOVERY.
2.7%; Score 66.6; DB 10; Length 58822; ar Local Similarity 88.9%; Pred. No. 2.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           // Match 2.7%; Score 66.6; DB 5; Length 110000;
Local Similarity 88.9%; Pred. No. 3.4e-06;
                                                                                                                                                                                                                                                              Score 66.6; DB 10; Length 58822;
Pred. No. 2.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 59748;
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ID ABZ71832 standard; CDNA; 452 BP.

DE Human cancer-associated gene SEQ ID NO 139.

PN WO20028399-A1.

PD 24-OCT-2002.

PA (TAKA-) TAKARA BIO INC.

Query Match

2.7%; Score 66.4; DB 8; Length 452; Best Local Similarity 79.0%; Pred. No. 1.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 86
                                                                                                                                                                                                                                             Query Match

Best Local Similarity 88.9%; Pred. No. 2.4e-06;
RESULT 1102

ID ABD33260 standard; DNA; 59748 BP.

DE Human cancer-associated (CA) gene HD07-044.

PA (SAGR-) SAGRES DISCOVERY INC.

Query Match

Best Local Similarity 88.9%; Pred. No. 2.4e-06;

RESULT 1103

ID ABD32848 standard; DNA; 104900 BP.

DE Human cancer-associated genomic DNA HD17-016.

PA (SAGR-) SAGRES DISCOVERY INC.

Query Match

DE Human cancer-associated genomic DNA HD17-016.

PA (SAGR-) SAGRES DISCOVERY INC.

Query Match

Best Local Similarity 88.9%; Pred. No. 3.3e-06;

RESULT 1104

ID AAR84800 standard; DNA; 110000 BP.

DE Nucleotide sequence of the human SPG4 gene.

PA (SAGR-) SAGRES DISCOVERY INC.

Query Match

PARR-SOURT 1104

PA (SAGR-) SAGRES DISCOVERY INC.

Query Match

PARR-SOURT 1104

PARR-SOURT 1104

PARR-SOURT 1104

PARR-SOURT 1104

PARR-SOURT 1104

PA (SAGR-) SAGRES DISCOVERY INC.

Query Match

PARR-SOURT 1104

PARR-SOURT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         O2-SEP-2004.
(SAGR-) SAGRES DISCOVERY INC.
ery Match
2.7%; Score 66.6; DB 13;
ery Match
--- cimilarity 84.3%; Pred. No. 4.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 66.4; DB 4;
ery Match 92.1%; Pred. No. 7.3e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAC12861 standard; cDNA; 163 BP.
Human secreted protein 5' EST, SEQ ID NO: 16936
EP1033401-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Manan cancer-associated genomic DNA HD17-001.
WO2004074320-A2.
                                                                                                                    ADE95788 standard; DNA; 58822 BP.
Human TCOF1 gene genomic DNA sequence.
W020039484-A2.
15-MAY-2003.
(SAGR-) SAGRES DISCOVERY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACH50650 standard; cDNA; 491 BP. Human mammary gland cDNA #55. US2003073623-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CNRS ) CNRS CENT NAT RECH SCI.
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2004.
-2004.
-fy Match
Best Local Similarity RESULT 1106
ID AAK87645 stand>
DE Human immur
PN W02001=
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(DRMA/) DRMANAC R T.
                                                           Best Local Similarity RESULT 1101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
RESULT 1107
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(GEST ) GENSET.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ouery Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query.
Beet Loca.
RESULT 1105
ID ABD328'
DE Humar
PN WO?
PD 0'
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PN

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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:29592
                              09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ETY Match 2.7%; Score 66.2; DB 4;
ery Match 89.9%; Pred. No. 1.1e-07;
                                                                                                                                                                                                     FISCHER C L.
ROSEN C A.
SOPPET D R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZENG Z.
LAFLEUR D W.
MOORE P A.
                                                                                                                                                                                                                                                                                                              ZENG Z.
LAFLEUR D W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PISCHER C L.
ROSEN C A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FISCHER C L.
ROSEN C A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZENG Z.
LAFLEUR D W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOPPET D R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OLSEN H.
EBNER R.
BIRSE C E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 1131
                                                                                                                                                                                                                                                                                                                                                                                      (OLSE/) OLSEN H.
(EBNE/) EBNER R.
(BIRS/) BIRSE C E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOPPET D R
RUBEN S M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OLSEN H.
EBNER R.
BIRSE C E.
                                                                                                                                                                                                                                                                                                                                                    MOORE P A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2004044191-A1.
                        WO200157182-A2.
                                                                                                                                                                                                                                                                             KYAW H.
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                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
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                                                                                                                                                                                                                                                                                                                                                    MOOR/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (BIRS/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EBNE/)
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                                                                                               Best
RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABZ76729 standard; DNA; 19969 BP.
Human lactadherin related secreted protein encoding DNA SEQ ID NO:3
                                                                                                                           Human musculoskeletal system related polynucleotide SEQ ID NO 2799, WO200155367-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                     ADJ30172 standard; DNA; 19183 BP.
Human musculoskeletal system-associated genomic DNA - SEQ ID 2799.
US2004009488-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-JAN-2001.
(REGC ) UNIV CALIFORNIA.
2.7%; Score 66.4; DB 4; Length 160552;
2r. ... Similarity 79.0%; Pred. No. 4.8e-06;
                                                                                                                                                                                                                                                     ABX59422 standard; cDNA; 19183 BP.
cDNA encoding novel human musculoskeletal system antigen #1766.
US2002147140-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (HUMA-) HUMAN GENOME SCI INC.
ry Match
L Local Similarity 79.0%; Pred. No. 1.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 35236;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 46404;
               09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 66.4; DB 4; Length 19183;
st Local Similarity 79.0%; Pred. No. 1.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                  2.7%; Score 66.4; DB 8; Length 19183; 79.0%; Pred. No. 1.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 2.7%; Score 66.4; DB 8; Length 19969;
Best Local Similarity 92.1%; Pred. No. 1.5e-06;
RESULT 1123
                                                                                                                                                                                                   Length 19183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAD02697 standard; DNA; 160552 BP.
Human glycosyl sulfotransferase-4 (GST-4) genomic DNA.
WO200106015-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ouery Match 2.7%; Score 66.4; DB 11;
Best Local Similarity 92.1%; Pred. No. 2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 11;
                                                                                                                                                                           HUMAN GENOME SCI INC.
2.7%; Score 66.4; DB 4;
Similarity 79.0%; Pred. No. 1.5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 2.7%; Score 66.4; DB 11
Best Local Similarity 79.0%; Pred. No. 2.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABD33462 standard; DNA; 196063 BP.
Human cancer-associated (CA) gene HD07-088.
WO2004058146-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACN44094 standard; DNA; 35236 BP. Human genomic sequence hCG23314. WO2003073826-A2.
                                                                                                          DNA; 19183 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACN44270 standard; DNA; 46404 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human genomic sequence hCG37971.
WO2003073826-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAK74780 standard; DNA; 138 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-SEP-2003.
(SAGR-) SAGRES DISCOVERY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-SEP-2003.
(SAGR-) SAGRES DISCOVERY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (APPL-) APPLERA CORP.
                                                                                                                                                                                                                                                                                                                         (ROSE/) ROSEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
ery Match
                                                                                                                                                                                                                 Local Similarity
                                                                                                          AAL36434 standard;
                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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RESULT 1126
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NO2003006485-A2.
WO200157182-A2.
                                                                                                                                                               02-AUG-2001
(HUMA-) HUM
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                                                       Query Match
                                                                                                                                                                                                   Match
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Human neuregulin gene single nucleotide polymorphism SNPBNRG680660.
WO200164876-A2.
07-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.7%; Score 66.2; DB 12; Length 146; 89.9%; Pred. No. 1.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.7%; Score 66.2; DB 12; Length 146; 89.9%; Pred. No. 1.1e-07;
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Length 138
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Query Match
                                                                            ADN41724 standard; DNA; 146 BP.
Novel human secreted protein polynucleotide seqid 846.
US2004044191-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADN41721 standard; DNA; 146 BP.
Novel human secreted protein polynucleotide segid 843.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel human secreted protein polynucleotide seqid 844. US2004044191-Al.
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07-NOV-2002
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Best Local S
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            RESULT 1141
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                                    LT 1132
AAK95773 standard; DNA; 401 BP.
Human neuregulin gene single nucleotide polymorphism SNP8NRG680652.
WO200164876-A2.
07-SEP-2001.
                                                                                                                                                                                                                                                                                                                  Best Local Similarity 72.2%; Pred. No. 1.9e-07;
RESULT 1133

ID AX897266 standard; DNA; 401 BP.
DE Human neuregulin gene single nucleotide polymorphism SNP8NRG680652.
PN W0200164817-A2.
PD 07-SSP-2001.
PA (DECO-) DECODE GENETICS EHF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAK§97267 standard; DNA; 401 BP.
Human neuregulin gene single nucleotide polymorphism SNP8NRG680660.
WO200164877-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-SEP-2001.
(DECO-) DECODE GENETICS EHF.
2.7%; Score 66.2; DB 4; Length 426;
2.7%; Score 66.2; DB 4; Length 426;
2.2%; Pred. No. 2e-07;
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Query Match 2.7%; Score 66.2; DB 4; Length 401;
Best Local Similarity 72.2%; Pred. No. 1.9e-07;
RESULT 1135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 401;
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DE Human neuregulin-1-associated gene 1 fragment SEQ ID NO: 573

PN US2002045577-A1.

PD 18-APR-2002.

PA (DECO-) DECODE GENETICS EHF.

Query March

2.7%; Score 66.2; DB 6; Length 401;

Best Local Similarity 72.2%; Pred. No. 1.9e-07;
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(DECO-) DECODE GENETICS EHF.
ery Match
2.7%; Score 66.2; DB 6; Length 401;
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(DECO-) DECODE GENETICS EHF.

2.7%; Score 66.2; DB 4; Length 401;

ery Match
72.2%; Pred. No. 1.9e-07;
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Human neuregulin-1-associated gene 1 fragment SEQ ID NO:
US2002045577-A1.
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(BECO) DECODE GENETICS EHF.

(CET) Match 2.7%; Score 66.2; DB 6;

ery Match 72.2%; Pred. No. 1.9e-07;
        72.2%; Pred. No. 1.9e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABT02037 standard; DNA; 401
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
RESULT 1138
ID ABT02036 standard; DN
DE Human neuregulin-1-as
PD 18-JUL-2002
PA (DECO-) DECODE GENETT
Best Local Similarity RESULT 1132
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Best Local Similarity
RESULT 1139
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(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-SEP-2001
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ABT01397 standard, DNA, 426 BP.
Human neuregulin-1-associated gene 1 fragment SEQ ID NO: 1426.
US2002045577-A1.
                                                                                                                                                                                                                                                                                                                                        _______ABTO02090 standard; DNA; 426 BP.
ABTO020990 standard; DNA; 426 BP.
US2002094954-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.7%; Score 66.2; DB 9; Length 471; 89.9%; Pred. No. 2.1e-07;
                                                                                                          Length 426;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 426;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human neuregulin gene insertion/deletion region #52. US2002165144-A1.
AAK98120 standard; DNA; 426 BP.
Human neuregulin gene insertion/deletion DNP8NRG49.
WO200164877-A2.
                                                                                                                                                                                                                                18-ASPR-2002.

18-ASPR-2002.

(DECO-) DECODE GENETICS EHF.

2.7%; Score 66.2; DB 6;

tery Match 12.2%; Pred. No. 2e-07;
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89.9%; Pred. No. 2.1e-07;
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Pred. No. 2.3e-07;
                                                      OPTION OF THE COLUMN OF T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH10353 standard; cDNA; 544 BP.
Human cDNA clone (3'-primer) SEQ ID NO:7188.
EP1074641742.
07-FEB-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OBECO-) DECODE GENETICS EHF ICELAND.
Query Match
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Human foetal brain cDNA #1388.
US2003073623-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACH26075 standard; cDNA; 471 BP. Human adult ovary cDNA #4455. US2003073623-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADH77652 standard; DNA; 426 BP.
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US2003073623-A1.
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(DECO-) DECODE GENETICS EHF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (DRWA) DRWANAC R T.
(LABA) LABAT I.
(STACY) STACHE-CRAIN B.
(DICK) DICKSON M C.
(JONE/) JONES L W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (DRWA) DRWANAC R T.
(LABA/) LABAT I.
(STAC/) STACHE-CRAIN B.
(DICK/) DICKSON M C.
(JONE/) JONES L W.
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(LABA/) LABAT I.
(STAC/) STACHE-CRAIN B.
(DICK/) DICKSON M C.
(JONE/) JONES L W.
                                                                                                          Query Match
Best Local Similarity
RESULT 1142
                                                                                                                                                                                                                                                                                                        Best Local Similarity
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RESULT 1144
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Length 1679;

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Humourassociated antigenic target (TAT) cDNA sequence #3626. WO2004060270-A2. CE-JUL-2004. GENENTROW
                                                                                                                                       Human tumour-associated antigenic target (TAT). cDNA sequence #2473.
WO2004060270-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADQ84271 standard; cDNA; 2186 BP.
Human tumour-associated antigenic target (TAT) cDNA sequence #1085.
WO2004060270-A2.
                                                                                                                                                                                                                                                                  2.7%; Score 66.2; DB 12; Length 2186; 89.9%; Pred. No. 5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.7%; Score 66.2; DB 12; Length 2186; 89.9%; Pred. No. 5e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 66.2; DB 4; Length 2368; Pred. No. 5.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
Query Match 2.7%; Score 66.2; DB 5; Length 4870;
Best Local Similarity 89.9%; Pred. No. 7.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.7%; Score 66.2; DB 10;
81.1%; Pred. No. 6.2e-07;
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89.9%; Pred. No. 5.7e-07;
           PA (HELI-) HELIX RES INST.

Query Match
2.7%; Score 66.2; DB 4;
Best Local Similarity 89.9%; Pred: No. 4.3e-07;
RESULT 1157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABV25122 standard; cDNA; 4870 BP.
Human prostate expression marker cDNA 25113.
WO200160860-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Leukaemia-related DNA sequence #2701.
WO2003039443-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 89.9%; Fred. N
RESULT 1160
LD AAH16324 standard; CDNA; 2368 BP.
DE Human CDNA sequence SEQ ID NO:15226.
PN EP1074617-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-MAY-2003.
(DEKR-) DEVI KREBSFORSCHUNGSZENTRUM.
(UYLU-) UNIV LUDWIG MAXIMILIANS.
(HAFE/) HAFERLACH T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADA53356 standard; cDNA; 3206 BP. Human coding sequence, SEQ ID 924 EP1293569-A2.
                                                                                                                ADQ85659 standard; cDNA; 2186 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HELI-) HELIX RES INST.
(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAK85164 standard; DNA; 5638 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADF82145 standard; DNA; 2788 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HELI-) HELIX RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-JUL-2004.
(GETH ) GENENTECH INC.
(WUTD/) WU T D.
(ZHOU/) ZHOU Y.
                                                                                                                                                                                                     (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
RESULT 1162
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Best Local Similarity
RESULT 1158
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SCHO/) SCHOCH C. (KERN/) KERN W.
                                                                                                                                                                                                                        (WUTD/) WU T D. (ZHOU/) ZHOU Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                      (WUTD/) WU T D.
(ZHOU/) ZHOU Y.
        07-FEB-2001
(HELI-) HEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABV76349 standard; cDNA; 1157 BP.
Human signal peptidase motif-containing protein 9.68-encoding cDNA.
CN1351026-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 2.7%; Score 66.2; DB 4; Length 1602; Local Similarity 89.9%; Pred. No. 4.2e-07;
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Query Match
Best Local Similarity 88.8%; Pred. No. 2.8e-07;
REGULT 1153
                            Length 544;
                                                                                                                                                                                                                                                                  Length 601;
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                                                                      ABX61812 standard; DNA; 601 BP.
Novel human transporter protein related polynucleotide #8.
022002142381-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABXXI810 standard; DNA; 601 BP.
Novel human transporter protein related polynucleotide #6
US2002142381-A1.
                                                                                                                                                                                                                                                                                                                                                  Novel human transporter protein related polynucleotide #7 US2002142381-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Single nucleotide polymorphism containing sequence #631
                            Score 66.2; DB 4;
Pred. No. 2.3e-07;
                                                                                                                                                                                                                                                              2.7%; Score 66.2; DB 8;
89.9%; Pred. No. 2.4e-07;
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Pred. No. 2.4e-07;
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Pred. No. 2.4e-07;
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Pred. No. 2.8e-07;
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ry Match
t Local Similarity 89.9%; Pred. No. 3.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAH06411 standard; cDNA; 763 BP.
Human cDNA clone (5'-primer) SEQ ID NO:3246.
EP1074617-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (WHED ) WHITEHEAD INST BIOMEDICAL RES. (ARFY.) AFFYMETRIX INC. 2.7%; Score 66. Local Similarity 87.7%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LT 1155
AAH18715 standard, CDNA, 1602 BP.
Human CDNA sequence SEQ ID NO:18986.
EP1074617-A2.
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ID AAH17499 standard; CDNA; 1679 BP.
BE Human CDNA sequence SEQ ID NO:16969.
PN EP1074617-A2.
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                                                                                                                                                                                                                                                                                                                              ABX61811 standard; DNA; 601 BP
                        Query Match 2.7%;
Best Local Similarity 89.9%;
RESULT 1149
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Local Similarity 89.9%;
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Local Similarity 89.9%;
(HELI-) HELIX RES INST.
                                                                                                                                                                        (GONG/) GONG F.
(KETC/) KETCHUM K A.
(DFRA/) DI FRANCESCO V.
(BEAS/) BEASLEY E M.
                                                                                                                                                                                                                                                                                                                                                                                                            (GONG/) GONG F.
(KETC/) KETCHUM K A.
(DFRA/) DI FRANCESCO V.
(BEAS/) BEASLEY E M.
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(KETC/) KETCHUM K A.
(DFRA/) DI FRANCESCO V.
(BEAS/) BEASLEY E M.
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                                                                                                                                                                                                                                                                                    Local Similarity
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Best Local Si
RESULT 1154
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Best Local Si
RESULT 1155
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Length 2788;

Length 3206;

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WO200159063-A2.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:41285.

WO200157182-A2.
09-AUG-2001.
09-AUG-2001.
2.7%; Score 66.2; DB 4; Length 13418;

Local Similarity 89.9%; Pred. No. 1.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAK82834 standard; DNA; 8789 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:37646.
WO200157182-A2.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:39976.
WO200157182-A2.
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Pred. No. 2e-06;
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Human testicular antigen encoding DNA fragment SEQ ID NO: 3133.
WO200155317-A2.
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                                                                                                                                                                                                                                                                                                                            Genomic sequence #473 encoding novel human enzyme polypeptide. WO200155301-A2. 02-AMG-2001. 02-AMG-2001. 2.7%: Score 66.2; DB 4; Length 7715; Et Local Similarity 89.9%; Pred. No. 1e-06;
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Human reproductive system related antigen DNA SEQ ID NO: 8605
WO200155320-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Genomic sequence #471 encoding novel human enzyme polypeptide WO200155301-A2.
                                                                                                                                      AUS42158 standard; DNA; 7712 BP.
Genomic sequence #474 encoding novel human enzyme polypeptide
WO200155301-A2.
                                                                                                                                                                                                             02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
2:7%; Score 66.2; DB 4; Length 7712;
st Local Similarity 89.9%; Pred. No. 1e-06;
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                                                                                      Length 5638,
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Human nervous system related polynucleotide SEQ ID NO 11904.
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(HUMA-) HUMAN GENOME SCI INC.
(ELY Match 2.7%; Score 66.2; DB 4;
(ery Match 89.9%; Pred. No. 1.1e-06;
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(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 66.2; DB 4;
ery Match 89.9%; Pred. No. 1.1e-06;
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(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 66.2; DB 4;
st Local Similarity 89.9%; Pred. No. 2.2e-06;
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                                                  09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 66.2; DB 4;
E. Local Similarity 89.9%; Pred. No. 8.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACN44886 standard; DNA; 26332 BP.
Human genomic sequence hCG39434.
W020030073826-A2.
12-SEP-2003.
(SAGR-) SAGRES DISCOVERY.
                                                                                                                                                                                                                                                                                                                DNA; 7715 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA; 9544 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ID ABL98481 standard; DNA; 32169
DB Human testicular antigen encod
PN W0200155317-A2.
PD '02-AUG-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
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Best Local Si
RESULT 1169
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Best Local Si
RESULT 1165
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AAK62257 standard; DNA; 33971 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:24069.
WO200157182-A2.
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Genomic DNA encoding novel human GABA transporter related protein.
US2002142381-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 54701;
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(SAGR-) SAGRES DISCOVERY INC.
Query Match
2.7%; Score 66.2; DB 13; Length 33362;
Best Local Similarity 89.9%; Pred. No. 2.3e-06;
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Query Match 2.7%; Score 66.2; DB 8; Length 63588;
                                                                                                                                                                                                                                                                                                                       NO: 3132
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Length 40645;
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                                                                                                                 AALO5916 standard; DNA; 32189 BP.
Human reproductive system related antigen DNA SEQ ID NO: 8604
WQ200155320-A2.
                                                                                                                                                                                                                             Length 32189;
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ID ABA19572 standard; DNA; 32189 BP.
DE Human nervous system related polynucleotide SEQ ID NO 11903
PN WO200159063-A2.
                                                                                                                                                                                                                                                                                                                  Human testicular antigen encoding DNA fragment SEQ ID W0200155317-A2.
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Best Local Similarity 89.9%; Pred. No. 3e-06;
RESULT 1182
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(HUMA-) HUMAN GENOME SCI INC.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 66.2; DB 5;
sery Match 2.2e-06;
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ry Match
t Local Similarity 89.9%; Pred. No. 2.3e-06;
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89.9%; Pred. No. 2.5e-06;
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                                                                                                                                                                                        02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
Query Match 2.7%; Score 66.2; DB 4;
Best Local Similarity 89.9%; Pred. No. 2.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                              2.7%; Score 66.2; DB 4;
89.9%; Pred. No. 2.2e-06;
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Human cancer-associated genomic DNA HD14-035.
WO2004074320-A2.
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Human genomic sequence hCG24071.
WO2003073826-A2.
                                                                                                                                                                                                                                                                                                    ABL98480 standard; DNA; 32189 BP
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(HUMA-) HUMAN GENOME SCI INC.
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(SAGR-) SAGRES DISCOVERY.
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(KETC/) KETCHUM K A.
(DFRA/) DI FRANCESCO V.
(BEAS/) BEASLEY E M.
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Best Local Similarity
RESULT 1181
                                                                      Best Local Similarity
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Best Local Similarity
RESULT 1177
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RESULT 1179
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81.1%; Pred. No. 4.8e-06;

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ADQ19501 standard; DNA; 135005 BP
 Best Local Similarity RESULT 1195
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Best Local Similarity 81.1%; Pred. No. 4.8e-06;
RESULT 1193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-DEC-2003.
(GEHO ) GEN HOSPITAL CORP.
2.7%; Score 66.2; DB 12; Length 128034;
                     LT 1183
AAD42934 standard; DNA; 70000 BP.
Human phospholipase A2 (PLA2), group VI (Ca2+-independent) gene.
US6410325-B1.
25-JUN-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 110000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 113079;
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                                                                                                                                                                                                                                                                                                                                                                      Length 110000;
                                                                                                                                                                                                                                                                Length 93443;
                                                                                                                                                                                                                                                                                                                   Length 110000;
                                                                                                                                                                                                                                                                                                                                                                                                                           Length 110000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 110000
                                                                                                                        Length 70000;
                                                                                                                                                                         ADQ19270 standard; DNA; 93443 BP.
Human soft tissue sarcoma-upregulated DNA - SEQ ID 2089
WO2004048938-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polymorphic human IDE genomic sequence, SEQ ID 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 66.2; DB 12;
Pred. No. 4.8e-06;
                                                                                                                                                                                                                                                                Score 66.2; DB 12;
Pred. No. 4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Match 2.7%; Score 66.2; DB 13;
Local Similarity 89.9%; Pred. No. 4.4e-06;
                                                                                                     PA (ISIS-) ISIS PHARM INC.
Query Match
Best Local Similarity 89.9%; Pred. No. 3.4e-06;
RESULT 1184
                                                                                                                                                                                                                                                                                                                                                                        Score 66.2; DB 4;
Pred. No. 4.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 66.2; DB 12;
Pred. No. 4.4e-06;
                                                                                                                                                                                                                                                                                                                   Score 66.2; DB 4;
Pred. No. 4.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                           Score 66.2; DB 6;
Pred. No. 4.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 66.2; DB 6;
Pred. No. 4.4e-06;
   89.9%; Pred. No. 3.2e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABD32998 standard; DNA; 113079 BP.
Human cancer-associated genomic DNA HD21-021.
02-58P-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human IDE gene variant DNA sequence SeqID187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADE41581 standard; DNA; 128034 BP.
Human IDE genomic sequence, SEQ ID 186.
WO2003054143+A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human IDE gene DNA sequence SeqID186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADE43582 standard; DNA; 128034 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA; 128034 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADH54060 standard; DNA; 128034 BP
                                                                                                                                                                                                                                            (PROT-) PROTEIN DESIGN LABS INC. ry Match 2.7%; Scolt Local Similarity 81.1%; Prec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SAGR-) SAGRES DISCOVERY INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                         2.78;
                                                                                                                                                                                                                                                                                                                   2.78;
                                                                                                                                                                                                                                                                                                                                                                        2.78;
                                                                                                                                                                                                                                                                                                                                                                                        72.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (NEUR-) NEUROGENETICS INC. (GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (NEUR-) NEUROGENETICS INC (GEHO ) GEN HOSPITAL CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GEHO ) GEN HOSPITAL CORP
Best Local Similarity RESULT 1183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADHS4059 standard;
                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity RESULT 1188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 1189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
RESULT 1190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2003054143-A2.
03-JUL-2003.
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                                                                                                                                                                                                                                 10-JUN-2004
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vuery match 2.7%; Score 66.2; DB 12; Length 135005; Best Local Similarity 89.9%; Pred. No. 4.9e-06; RESULT 1196
                                                                                                                                                                ADPES634 standard; DNA; 137908 BP.
Human sequence from clone 914P14 on chromosome Xq23 Contains DNA
WO2003072827-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-FEED-2003.
(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
2.7%; Score 66.2; DB 10; Length 177531;
ery Match
2.7%; Score 66.2; DB 10; Length 177531;
                                                                                                                                                                                                                                                                   (CHIL-) CHILDREN'S HOSPITAL MEDICAL CENT.

ry Match 2.7%; Score 66.2; DB 11; Length 137908; t Local Similarity 85.1%; Pred. No. 5e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 2.7%; Score 66.2; DB 10; Length 161280;
Best Local Similarity. 81.1%; Pred. No. 5.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.7%; Score 66.2; DB 10; Length 161280; 81.1%; Pred. No. 5.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.7%; Score 66.2; DB 10; Length 177531; 81.1%; Pred. No. 5.7e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2.7%; Score 66.2; DB 10; Length 177531;
81.1%; Pred. No. 5.7e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 177531;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 177531;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACF62732 standard; DNA; 177531 BP. Cancer based on CYP3A5 related polynucleotide SEQ ID NO:660 WO2003013534-A2.
Human soft tissue sarcoma-upregulated DNA - SEQ ID 2320 WO2004048938-A2.
                                                                                                                                                                                                                                                                                                                                                               ADD67054 standard; cDNA; 161280 BP.
Human lung tumour-specific related cDNA, SEQ ID No 746.
WO200292001-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MRP1 based cancer related nucleic acid SEQ ID NO:660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 2.7%; Score 66.2; DB 8; Best Local Similarity 81.1%; Pred. No. 5.7e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.7%; Score 66.2; DB 8;
81.1%; Pred. No. 5.7e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA; 177531 BP.
DNA sequence SEQ ID NO:660.
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DNA sequence SEQ ID NO:660.
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Human UGT1A1 gene sequence SEQ ID NO:660.
WO2003013536-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
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(EPID-) EPIDAUROS BIOTECHNOLOGIE AG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human lung tumour antigen cDNA #640. US2003118599-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                            Local Similarity
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ID ADB96919 standard;
DE Human MDR1 related
PN WO2003013537-A2.
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RESULT 1213
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(SAGR-) SAGRES DISCOVERY INC.

Query Match | Sagre 66.2; DB 13; Length 310268;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 210204;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.7%; Score 66.2; DB 12; Length 254396; 89.9%; Pred. No. 7e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.7%; Score 66.2; DB 11; Length 304905; 89.9%; Pred. No. 7.7e-06;
                                                                                                                 Length 177531;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 202100;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADQ18927 standard; DNA; 210204 BP.
Human soft Lissue sarcoma-upregulated DNA - SEQ ID 1746.
WO2004048938-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7 1209
ADL18884 standard, DNA; 247509 BP.
Recoarthitis associated polymorphic nucleotide #216.
WO2003054166-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADQ97557 standard; DNA; 254396 BP.
Human cancer associated sequence HD09-013, SEQ ID 534.
WO2004060304-A2.
                                                                                                                                                                                                 Osteoarthritis-associated polymorphic nucleotide #44.
WO2003054166-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOCNULATIONS.

(INCY-) INCYTE GENOMICS INC.

(INCY-) INCYTE GENOMICS INC.

2.7%; Score 66.2; DB 10;

ery Match

2.7%; Pred. No. 6.9e-06;
                                                                        31-JUL-2003.
(SJUD-) ST JUDE CHILDREN'S RES HOSPITAL.
ery Match 2.7%; Score 66.2; DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 66.2; DB 12;
Pred. No. 6.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 12;
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(PROT-) PROTEIN DESIGN LABS INC.

ery Match 2.7%; Score 66.2; DB 12

Bt Local Similarity 81.1%; Pred. No. 6.3e-06;
                                                                                                                                                                                                                                                                                                                         ADE43315 standard; DNA; 202100 BP.
Human IDE/ KNSL1 genomic sequence, SEQ ID 484.
WO2003054143-A2.
                  ADH74617 standard; DNA; 177531 BP.
Human BAC clone GS1-259H13 CYP3A5 genomic DNA.
US2003143537-A1.
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Human cancer-associated genomic DNA HD14-033.
WO2004074320-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADH54357 standard; DNA; 202100 BP.
Human IDE/KNSL1 gene DNA sequence SeqID484.
US2003224380-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-APR-2003.
(GENO-) GENOME THERAPEUTICS CORP.
                                                                                                                                                                             ADL13512 standard; DNA; 178870 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADP75180 standard; DNA; 304905 BP
Human Endophilin 2 gene.
WO2003031594-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.7%;
81.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-DEC-2003.
(GEHO ) GEN HOSPITAL CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                   (NEUR-) NEUROGENETICS INC. (GEHO ) GEN HOSPITAL CORP.
                                                                                                                           Best Local Similarity
RESULT 1205
ID ' ADL13512 standard; DN
DE Osteoarthritis-associ
PN WO2003054166-A2.
PD 03-JUL-2003.
PA (INCY-) INCYTE GENOMI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
RESULT 1208
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Best Local Similarity
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                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 1206
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                                                                                                                     Query Match
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RESULT 1204
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Human cardiovascular system associated polypeptide-related DNA SeqID1599.
US2004005575-A1.
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WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAK66109 standard, DNA, 11684 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:20921.
                                                           Human cardiovascular system antigen genomic DNA SEQ ID No 1599 WO200155321-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 17792;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 17792;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1218
TD AAS3727 standard; DNA; 17792 BP.
DE Human genomic DNA for novel endocrine antigen, SEQ ID No 681.
PN WO200155319-A2.
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(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 66; DB 4; Length 20245;
ery Match 2.7%; Pred. No. 1.9e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           y Match 2.7%; Score 66; DB 4; Length 17792;
Local Similarity 79.6%; Pred. No. 1.8e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 11684;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 17792;
                                                                                                                                                                                                                                                                               DB 4; Length 2076;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 3685;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human cardiovascular system related genomic DNA #359 US2003059908-A1.
                                                                                                                                                                                                                                                                                                                                            cDNA; 3685 BP.
human diagnostic protein #9529.
                                                                                                                                                                                                                                                         (HELI-) HELIX RES INST.

2.7%; Score 66; DB 4; 1
Best Local Similarity 72.1%; Pred. No. 5.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.7%; Score 66; DB 5; 1
87.8%; Pred. No. 7.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.7%; Score 66; DB 4; I
93.2%; Pred. No. 1.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HUMA-) HUMAN GENOME SCI INC.

2.7%; Score 66; DB 4; I

2.0cal Similarity 79.6%; Pred. No. 1.8e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.7%; Score 66; DB 10;
79.6%; Pred. No. 1.8e-06;
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ABD33100 standard; DNA; 313287 BP.
Human cancer-associated (CA) gene HD07-008.
WO2004058146-A2.
                                                                                                                                                              AAH16545 standard; CDNA; 2076 BP.
Human CDNA sequence SEQ ID NO:15602.
EP1074617-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAS36099 standard; DNA; 17792 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T 1219
ADE46793 standard; DNA; 17792 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADJ08211 standard; DNA; 17792 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAK72318 standard; DNA; 20245 BP
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(HUMA-) HUMAN GENOME SCI INC.
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(HUMA-) HUMAN GENOME SCI INC.
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                                                                                                     Query Match
Best Local Similarity
RESULT 1214
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Best Local Similarity
RESULT 1222
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Best Local Similarity
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                          AAS73725 standard;
DNA encoding novel
WO200175067-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 1216
                                                                                                                                                                                                                                                                                                                                                                                                          11-OCT-2001.
(HYSE-) HYSEQ INC.
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ACC45361 standard; DNA; 26928 BP.
Human HBM gene fragment #2.
WO200292764-A2.
                                                                      21-NOV-2002.
(GENO-) GENOME THERAPEUTICS CORP.
(AMHP ) WYETH.
                                                                                                                                              Best Local Similarity RESULT 1232
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RESULT 1237
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RESULT 1239
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Best Local Similarity
RESULT 1233
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09-AUG-2001.
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Best Local S
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                PA PA
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Secreted protein gene 12 genomic fragment HAUA183, SEQ ID NO:592.
WO200276488-A1.
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ry Match 2.7%; Score 66; DB 10; Length 20245;
t Local Similarity 87.8%; Pred. No. 1.9e-06;
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Query Match
Best Local Similarity 87.8%; Pred. No. 1.9e-06;
                                                                                                               Length 20245;
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(UVCR-) UNIV CREIGHTON SCHOOL MEDICINE.

2, 7%; Score 66; DB 6; Length 26928;

Query Match

Best Local Similarity 76.4%; Pred. No. 2.3e-06;
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ADAS7727 standard, DNA; 20245 BP.
BAC frament containing human secreted protein gene #302.
27-DEC-2002.
ABK69845 standard; DNA; 20245 BP.
Human secreted protein gene 22 genomic DNA fragment
WO200226931-A2.
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Human high bone mass (HBM) polynucleotide clone #2.
MC0200192891-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADC74613 standard, DNA; 20245 BP.
Human secreted protein-related DNA - SEQ ID 1246.
WO2003038063-A2.
                                                                                                                                                      ACCOURGO standard; CDNA; 20245 BP.
Human secreted protein BAC clone SEQ ID NO 1040.
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(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 66; DB 10;
ery Match 7.1%; Pred. No. 1.9e-06;
                                                                        04-APR-2002.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 66; DB 6; Tt Local Similarity 87.8%; Pred. No. 1.9e-06;
                                                                                                                                                                                                                            28-NOV-2002.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 66; DB 8;
st Local Similarity 87.8%; Pred. No. 1.9e-06;
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2.7%; Score 66; DB 5; 1
Best Local Similarity 76.4%; Pred. No. 2.3e-06;
RESULT 1230
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Human secreted protein related DNA #SEQ ID 784,
WO2003004522-A2.
16-JAN-2003.
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ABAB2620 standard; DNA; 26928 BP.
Human HBM gene region b527d12-h_contig306G.
W0200177327-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADD38133 standard; cDNA; 20245 BP. cDNA clone in ArCC deposit #27. WO200290526-A2.
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Best Local Si
RESULT 1227
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RESULT 1231
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Query Match
Best Local Similarity 76.4%; Pred. No. 2.3e-06;
RESULT 1236
HEBOUT 1236
Human neuroblastoma cell line NB-1 1p36 nucleotide sequence SEQ ID NO:60.

PN WOO20116311-A1.
PD 08-MAR-2001.
PD PARP 2001.
PD PARP 2001.
PD PARP 2001.
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Genomic sequence #465 encoding novel human connective tissue polypeptide.
WO200155341-A1.
02-AUG-2001.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:40693.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:40695.
WO200157182-A2.
                                                                                                                                                                                                                                           Length 26928;
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Length 26928;
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BAC clone containing segments of the human Zmax1 gene #2.
US6780609-B1.
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BAC clone containing segments of the human Zmax1 gene #2.
US2004176582-A1.
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2.7%; Score 65.8; DB 4;
Similarity 90.9%; Pred. No. 1.8e-07;
  2.7%; Score 66; DB 8; 176.4%; Pred. No. 2.3e-06;
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Query Match 2.7%; Score 66; DB 13;
Best Local Similarity 76.4%; Pred. No. 2.3e-06;
RESULT 1235
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(CHIB-) CHIBA PREPECTURE.
2.77; Score 66; DB 5; I
t Local Similarity 72.1%; Pred. No. 3.3e-06;
                                                                                                                                                                                                                                             2.7%; Score 66; DB 10; 76.4%; Pred. No. 2.3e-06;
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                                                                                                                                                                                                                                                                                                                         ADE82430 standard; DNA; 26928 BP.
Human DNA sequence related to the invention #2.
WO200292015-A2.
                                                                              ADB98061 standard; DNA; 26928 BP.
HBM-related clone contig b527d12-h contig306G.
W0200292000-A2.
21-NOV-2002.
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AAK64739 standard; DNA; 58708 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:19551.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human immune/haematopoietic antigen genomic sequence SEQ ID NO:22023. WO200157182-A2.
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(SAGR-) SAGRES DISCOVERY INC.
2.7%; Score 65.8; DB 13; Length 32229;
Et Local Similarity 90.9%; Pred. No. 2.9e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.7%; Score 65.8; DB 5; Length 24183; 90.9%; Pred. No. 2.4e-06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 12026;
                                                             2.7%; Score 65.8; DB 5; Length 1867; 90.9%; Pred. No. 5.9e-07;
                                                                                                                                                                                                                                          Length 1887;
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Human cancer associated gene genomic sequence SEQ ID NO:58.
WO2004074321-A2.
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90.9%; Pred. No. 2.8e-06;
                                                                                                                                                                                                                                             2.7%; Score 65.8; DB 4; 90.9%; Pred. No. 5.9e-07;
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(HUMA-) HUMAN GENOME SCI INC.
Query Match
2.7%; Score 65.8; DB 4;
Best Local Similarity 90.9%; Pred. No. 1.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        / Match 2.7%; Score 65.8; DB 4;
Local Similarity 90.9%; Pred. No. 1.7e-06;
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Human cancer-associated (CA) gene HD07-104.
WO2004058146-A2.
                                                                                                                          Al61210 standard; cDNA; 1887 BP.
Human polynucleotide SEQ ID NO 5199.
WC200153312-Al.
26-JUL-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-JUL-2001.
(UYAL-) UNIV ALLEGHENY HEALTH SCI.
(UYJE-) UNIV JEFFERSON THOWAS.
(UYOU-) UNIV OULU.
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Human genomic sequence hCG34092.
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Human gene for collagen COL9A1.
US6265157-B1.
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RESULT 1253
ID ACM43982 standard; DNA; 24898 BP.
DE Human genomic sequence hCG23625.
PN WO2003073826-A2.
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ID AAK67211 standard; DNA; 12026 BP
              (UYOU-) UNIV JEFFERSON THOMAS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
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(SAGR-) SAGRES DISCOVERY INC.
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                                                                           Best Local Similarity
RESULT 1249
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Best Local Similarity
RESULT 1250
                                                                                                                                                                                                                     (HYSE-) HYSEQ INC.
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                                                                                                       ABK42567 standard; DNA; 290 BP.
Genomic sequence #466 encoding novel human connective tissue polypeptide.
WO200155343-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genomic sequence #467 encoding novel human connective tissue polypeptide. WO200155343-A1.
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Genomic sequence #468 encoding novel human connective tissue polypeptide.
NO200155343-Al.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:38778.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match
ery Match
st Local Similarity 85.9%; Pred. No. 3.9e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PD 02-AUG-2001.

PA (HUMA-) HUMAN GENOME SCI INC.

Query Match 2.7%; Score 65.8; DB 4; Length 291;

Best Local Similarity 85.9%; Pred. No. 2.1e-07;

RESULT 1245
                                                                                                                                                             PD 02-AUG-2001.

PA (HUMA-) HUMAN GENOME SCI INC.

Query March

Best Local Similarity 85.9%; Pred. No. 2.1e-07;

RESULT 1241
                                                                                                                                                                                                                                                                                                                                                                                                        Length 290;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 291;
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match 2.7%; Score 65.8; DB 4; Length 290;
Best Local Similarity 85.9%; Pred. No. 2.1e-07;
RESULT 1240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 290;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 2.7%; Score 65.8; DB 4; Length 291; Local Similarity 85.9%; Pred. No. 2.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 291;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-MAR-2003.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 65.8; DB 9;
2.1. Local Similarity 85.9%; Pred. No. 2.1e-07;
17 1248
                                                                                                                                                                                                                                                                                                                                                                           (HUMA-) HUMAN GENOME SCI INC.

2.7%; Score 65.8; DB 9;
t Local Similarity 85.9%; Pred. No. 2.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-MAR-2003.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 65.8; DB 9;
t Local Similarity 85.9%; Pred. No. 2.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HUMA-) HUMAN GENOME SCI INC.

2.7%; Score 65.8; DB 9;
t Local Similarity 85.9%; Pred. No. 2.1e-07;
                                                                                                                                                                                                                                                                                     ADB60723 standard; DNA; 290 BP.
Connective tissue related genomic DNA #466.
US2003054375-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADB60722 standard; DNA; 290 BP.
Connective tissue related genomic DNA #465.
US2003054375-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADB60724 standard; DNA; 291 BP.
Connective tissue related genomic DNA #467.
US2003054375-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Connective tissue related genomic DNA #468.
US2003054375-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAS21879 standard; DNA; 1867 BP.
Human collagen gene COL9A1 5' UTR.
USCS5157-B1.
24-UL-2001.
(UYAL-) UNIV ALLEGHENY HEALTH SCI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    T 1244
ABK42568 standard; DNA; 291 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADB60725 standard; DNA; 291 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (HUMA-) HUMAN GENOME SCI INC.
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Best Local Si
RESULT 1243
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Best Local Si
RESULT 1246
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Best Local Si
RESULT 1242
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Best L RESULT 1 ID ABK DE GEN PN WO2 PD 02-

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AAK85587 standard; DNA; 132 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:40399.
WO200157182-A2.
                                                                                                                                                                                                                                    AAK85588 standard; DNA; 132 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:40400.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                          TARK80586 standard; DNA; 142 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:35398.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human digestive system antigen genomic sequence SEQ ID NO: 3986 WO200155314-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABL98078 standard; DNA; 152 BP.
Human testicular antigen encoding DNA fragment SEQ ID NO: 2730.
WQ200155317-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AALOS193 standard; DNA; 152 BP.
Human reproductive system related antigen DNA SEQ ID NO: 7881.
WO200155320-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ID No 482
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oz-Audra-) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 4; Length 152;
rr Local Similarity 88.8%; Pred. No. 1.7e-07;
                                                                                                                                                                                                                                                                                                                                                            Length 132;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABA17343 standard; DNA; 151 BP.
Human nervous system related polynucleotide SEQ ID NO 9674.
WO200159063-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAS12528 standard, DNA; 145 BP.
Human genomic DNA for novel endocrine antigen, SEQ
WO200155319-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAS33446 standard; DNA; 152 BP.
DNA encoding human secreted protein, Seg ID No 729.
WO200155326-A2.
                                                                                                                09-AUG-2001.
(HUWA-) HUMAN GENOME SCI INC.
(HUWA-) HUMAN GENOME SCI INC.
(ery Match 2.7%; Score 65.6; DB 4;
ery Match 1.5e-07;
                                                                                                                                                                                                                                                                                                                               (HUMA-) HUMAN GENOME SCI INC.
ry Match
t Local Similarity 88.8%; Pred. No. 1.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HUMA-) HUMAN GENOME SCI INC.
ry Match
t Local Similarity 88.8%; Pred. No. 1.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HUMA-) HUMAN GENOME SCI INC.

2.7%; Score 65.6; DB 5;

L Local Similarity 88.8%; Pred. No. 1.7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HUMA-) HUMAN GENOME SCI INC.
ry Match
t Local Similarity 88.8%; Pred. No. 1.7e-07;
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ry Match
t Local Similarity 88.8%; Pred. No. 1.7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (HUMA-) HUMAN GENOME SCI INC.

2.7%; Score 65.6; DB 4;

t Local Similarity 88.8%; Pred. No. 1.6e-07;
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No. 1.7e-07;
    88.8%; Pred. No. 1.5e-07;
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2.7%; Score
Similarity 88.8%; Pred.
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                                                                                                                                                                                   Best Local Similarity RESULT 1268
    Best Local Similarity
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RESULT 1270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                             09-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AASO9301 standard; DNA; 319608 BP.
Human schizophrenia associated gene g35030 and biallelic markers Al-A71.
ery Match
2.7%; Score 65.8; DB 5; Length 319608;
st Local Similarity 85.9%; Pred. No. 1e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAK67355 standard; DNA; 109 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:22167.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAK67159 standard; DNA; 109 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:21971.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAK86351 standard; DNA; 128 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:41163.
WO200157182-A2.
09-AUG-2001.
                                                                       2.7%; Score 65.8; DB 11; Length 217409;
30.9%; Pred. No. 8.2e-06;
                                                                                                                                                                                                                                                            Length 254396;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.7%; Score 65.8; DB 3; Length 319608; 85.9%; Pred. No. 1e-05;
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2.7%; Score 65.6; DB 4; Length 128;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human nervous system related polynucleotide SEQ ID NO 7610.
WO200159063-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 92;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADAS7641 standard; DNA; 101 BP.
BAC fragment containing human secreted protein gene #49.
W02002102994-A2.
27-DEC-2002.
(HUMA-) HUWAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                            AAHS1601 standard; DNA; 319608 BP.
Human chromosome 13q31-q33 genomic nucleotide sequence.
WO200058510-A2.
                                                                                                                   AUQ97557 standard; DNA; 254396 BP.
Human cancer associated sequence HD09-013, SEQ ID 534.
WO2004060304-A2.
                                                                                                                                                                                                                                                               12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.7%; Score 65.6; DB 10;
88.8%; Pred. No. 1.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 65.6; DB 4;
ery Match 2.7%; Pred. No. 1.4e-07;
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09-AUG-2001.
(HUMAA) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 4;
ery Match
2.7%; Score 65.6; DB 4;
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88.8%; Pred. No. 1.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Match 2.7%; Score 65.6; DB 5; Local Similarity 88.8%; Pred. No. 1.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AACIS312 standard, cDNA; 124 BP.
Human secreted protein 5' EST, SEQ ID NO: 19387
                                                                                                                                                                                                    Nac.-2004.
(SAGR-) SAGRES DISCOVERY INC.
(ST. SAGRES DISCOVERY INC.
(ery Match 2.7%; Score 65.8; DB 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA; 92 BP
                                                                                             Local Similarity 90.9%;
                        12-SEP-2003.
(SAGR-) SAGRES DISCOVERY.
                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABA15279 standard;
WO2003073826-A2.
                                                                                                                                                                                                                                                                                                                                                                                                   05-OCT-2000.
(GEST ) GENSET.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EP1033401-A2.
06-SEP-2000.
                                                                       Query Match
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Best 1 RESULT 1

Query | Best L RESULT 1

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ADJ08078 standard; DNA; 290 BP.
Human cardiovascular system associated polypeptide-related DNA SeqID1466.
                                                                                                                                                                                                           ARK70645 standard; DNA; 278 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:25457.
WO200157182-A2.
                                         AAK74855 standard; DNA; 265 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:29667.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human cardiovascular system antigen genomic DNA SEQ ID No 1466.
WO200155321-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-WAR-2003.
(HUWA-) HUMAN GENOME SCI INC.
Lery Match 2.7%; Score 65.6; DB 10; Length 290;
Lery Match 1 cimilarity 88.8%; Pred. No. 2.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CALB-) CALIFORNIA INST BIOLOGICAL RES.

ry Match
2.7%; Score 65.6; DB 2; Length 283;
t Local Similarity 88.8%; Pred. No. 2.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.7%; Score 65.6; DB 12; Length 283; 88.8%; Pred. No. 2.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OS-JAN-2004.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 65.6; DB 13; Length 290;
ery Match anilarity 88.8%; Pred. No. 2.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.7%; Score 65.6; DB 5; Length 302;
88.8%; Pred. No. 2.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ама18165 standard; DNA; 302 BP.
Human nervous system related polynucleotide SEQ ID NO 10496.
WQZ00159063-AZ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADE46660 standard; DNA; 290 BP.
Human cardiovascular system related genomic DNA #226.
US2003059908-A1.
                                                                                                                                                                                                                                                                                                                                                                                                   AAG63862 standard; cDNA; 283 BP.
AP2 sequence obtd. by PCR for tumour specific DNA 209411531-A1.
26-MAY-11994.
                                                                                                     09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
(ELY Match 2.7%; Score 65.6; DB 4;
Lery Match 88.8%; Pred. No. 2.3e-07;
                                                                                                                                                                                                                                                                                  09-ACG-2001.
(HUMA-) HUMAN GENOME SCI INC.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 4;
(ery Match 2.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.7%; Score 65.6; DB 4;
88.8%; Pred. No. 2.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 65.6; DB 9;
Pred. No. 2.3e-07;
  88.8%; Pred. No. 2.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AD004061 standard; DNA; 283 BP.
Human AP-PCR gene fragment (APdelta2) #2.
US2004091889-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADB97627 standard; DNA; 283 BP. APdelta2 DNA fragment #2. US6566053-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAS35966 standard; DNA; 290 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
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88.8%;
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RESULT 1293
ID ABA18165 standard; D
  Best Local Similarity
                                                                                                                                                                        Best_Local Similarity
RESULT 1286
                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity RESULT 1287
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(STRA-) STRATAGENE.
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(STRA-) STRATAGENE.
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Best Local Similarity
RESULT 1292
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   лик/U933 standard; DNA; 187 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:25745.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADI97188 standard; DNA; 251 BP.
Human respiratory system associated polypeptide-related DNA SeqID652.
US2003077704-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAKČEGSG standard; DNA; 262 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:20448.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                             AAK67885 standard, DNA; 156 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:22697.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAS28218 standard; DNA; 251 BP.
Genomic sequence #58 encoding for novel human respiratory antigen.
WO200155448-A1.
                    AAS39966 standard; DNA; 152 BP.
Genomic sequence #385 encoding human colon associated polypeptide.
WO200155302-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 251;
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(HUMA) HUMAN GENOME SCI INC.
12.7%; Score 65.6; DB 4; Length 187;
Fr Local Similarity 88.8%; Pred. No. 1.9e-07;
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2.7%; Score 65.6; DB 4; Length 262;
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                                                                                                                                                                                               ADB12926 standard; DNA; 152 BP.
Human novel colon related polypeptide DNA SEQ ID NO 863.
US2003050221-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human respiratory system associated genomic DNA seq id US2003215893-A1.
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(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 10;
st Local Similarity 88.8%; Pred. No. 2.2e-07;
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(HUMA-) HUMAN GENOME SCI INC.
(HVA-) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 11;
ery Match 2.2e-07;
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02-MCJ-2001.

(HUANG-) HUMAN GENOME SCI INC.

2.7%; Score 65.6; DB 4;

ery Match

2.7%; Score 65.6; DB 4;

2.7%; Pred. No. 2.2e-07;
                                                                              02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
(Ery Match 2.7%; Score 65.6; DB 5;
ery Match 88.8%; Pred. No. 1.7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (HUMA-) HUMAN GENOME SCI INC.
ry Match
L Local Similarity 88.8%; Pred. No. 1.7e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAC21709 standard; cDNA; 180 BP.
Human secreted protein 5' EST, SEQ ID NO: 25784
EP1033401-A2.
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RESULT 1278
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                                                                                                                                                                                                                                                                                         (RUBE/) RUBEN C A.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GEST ) GENSET.
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Best Local Si
RESULT 1283
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RESULT 1276
ID AAS3996
DE Genomic
PN WO200159
PD 02-AUG-?
PA (HUMA-)
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Best I RESULT 1

PERGE

Length 283;

Length 278,

Length 265

Length 290;

us-10-002-967a-58.rng.spdi

Query Match

Query Match

Match

(HUMA-)

Match

Query

Query Match

Query Match

Match

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AAK77558 standard; DNA; 309 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:32370.
WO200157182-A2.
                   2.7%; Score 65.6; DB 11; Length 308; 88.8%; Pred. No. 2.5e-07;
                                                                                                                                                                                                                                                                     Score 65.6; DB 12; Length 308;
Pred. No. 2.5e-07;
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ABA19007 standard; DNA; 321 BP.
HMAIN nervous system related polynucleotide SEQ ID NO 11338
WO200159063-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Abaleoo standard; DNA; 321 BP.
Human nervous system related polynucleotide SEQ ID NO 11339.
WO200159063-A2.
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Human nervous system related polynucleotide SEQ ID NO 11337.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABA14684 standard; DNA; 321 BP.
Human nervous system related polynucleotide SEQ ID NO 7015.
WQ200159063-A2.
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ry Match
L Local Similarity 88:8%; Pred. No. 2.5e-07;
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88.8%; Pred. No. 2.5e-07;
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2.7%; Score 65.6; DB 5;
Local Similarity 88.8%; Pred. No. 2.5e-07;
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                                                                       ADE44040 standard; cDNA; 308 BP.

Human cDNA associated with breast cancer #87.
US2003104366-A1.
05-JUN-2003.
(DILL/) JIANG Y.
(MITC.) MITCHAM J L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-SEP-2001.
(MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     . (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC. Query Match 2.7%; Score 65.6: D
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Human ovarian cancer DNA marker #12246.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADI73224 standard; DNA; 316 BP.
Human ovarian cancer DNA marker #5966
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(HUMA-) HUMAN GENOME SCI INC.
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                                                                                                                                                                                                                                                                         2.7%;
Best Local Similarity 88.8%;
RESULT 1304
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(HARL/) HARLOCKER S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ouery Match
Best Local Similarity
RESULT 1306
                                  Best Local Similarity
RESULT 1303
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RESULT 1305
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                                                                                                                                                                                                                                                                                                              2.7%; Score 65.6; DB 4; Length 308; 88.8%; Pred. No. 2.5e-07;
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                                                                                                                                           Length 307;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 308;
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Human nervous system related polynucleotide SEQ ID NO 10429.
WO200159063-A2.
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Human ovarian cancer related cDNA clone SEQ ID NO:316. W0200195581-A2. G6-DEC-2001. (CORI-) CORIXA CORP.
                                                                                                                                                                                                                         Human breast cancer associated 13053 coding sequence. WO200060076-A2.
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Human breast cancer-associated polypeptide cDNA #87.
US2003166022-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.7%; Score 65.6; DB 4;
88.8%; Pred. No. 2.5e-07;
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Human tumour-related DNA sequence - SEQ ID No 88
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2.7%; Score 65.6; DB 5;
Similarity 88.8%; Pred. No. 2.5e-07;
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Pred. No. 2.5e-07;
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88.8%; Pred. No. 2.5e-07;
                                                                                                                                         2.7%; Score 65.6; DB 6;
88.8%; Pred. No. 2.4e-07;
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US2002102602-A1.
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Human breast tumour polynucleotide #87.
US2002085998-Al.
                                                                                                                                                                                                                                                                                                                                                                         AAS47033 standard; cDNA; 308 BP.
Human breast cancer cDNA clone 13053.
WO200179286-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABK95068 standard; cDNA; 308 BP. Human breast tumour cDNA 13053. US6387697-B1.
                                                                                                                                                                                                        AAF17603 standard; cDNA; 308 BP
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88.8%;
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(DILL/) DILLON D C.
(MITC/) MITCHAM J L.
(XUJ/) XU J.
             RESULT 1294

ID ABL77338 standard; c
DE Human ovarian cancer
PN WC200192581-A2.
PD 06-DEC-2001.
PA (CORI-) CORIXA CORP.
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                                                                                                                                                        Best Local Similarity
RESULT 1295
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RESULT 1296
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RESULT 1301
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                                                                                                                                                                                                                                                                   12-OCT-2000
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Length 316;

Length 309

Length 321

Length 321

Length 321

Length 321

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(HUMA-) HUMAN GENOME SCI INC.
Query Match 2.7%;
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02-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAK65488 standard; DNA; 340 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:20300.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAK65487 standard; DNA; 340 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:20299.
WO200157182-A2.
                                                                                                                ABZ73926 standard; DNA; 336 BP.
Secreted protein gene 105 genomic fragment HE8SG96, SEQ ID NO:1073.
WO200277013-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AASSTGO standard; DNA; 347 BP.
Human cardiovascular system antigen genomic DNA SEQ ID No 1260.
WO200155321-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAS35765 standard; DNA; 347 BP.
Human cardiovascular system antigen genomic DNA SEQ ID No 1265.
WO200155321-A2.
                                                                                                                                                                                                                                                                                                                                                                       Length 336;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 336;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.7%; Score 65.6; DB 6; Length 337; 88.8%; Pred. No. 2.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          vuery Match 2.7%; Score 65.6; DB 4; Length 340; Best Local Similarity 88.8%; Pred. No. 2.6e-07; RESULT 1318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 347;
                                                         Length 336;
                                                                                                                                                                                                                Length 336;
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                                                                                                                                                                                                                                                                                                                                                                                                                              ABZ67514 standard; DNA; 336 BP.
Human secreted protein encoding genomic DNA SEQ ID NO 1037.
WO200277186-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABL77353 standard; cDNA; 337 BP.
Human ovarian cancer related cDNA clone SEQ ID NO:331.
WO200192581-A2.
                                                                                                                                                                                                                        Love 292787-A2.

Full 21-VOV-2002.

PA (HUMA-) HUMAN GENOME SCI INC.

Query Match

Best Local Similarity 88.8%; Pred. No. 2.6e-07;

RESULT 1314

ID ABZ67514 standard; DNA; 336 BP.

DE Human secreted protein encodimans accreted protein encodimans.

PD 03-OCT-200?
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LUCATO PA (HUMA-) HUMAN GENOME SCI INC.

QUETY MATCh

BESTLE LOCAL Similarity 88.8%; Pred. No. 2.6e-07;

RESULT 1315

ID ABL77353 standard; CDNA; 337 BP.

DE Human ovarian cancer related —

PN WC200192581-A2.

PA (CCT.
                                                                                                                                                                 PD 03-CCT-2002.

PA (HUMA-) HUMAN GENOME SCI INC.

Query Match

2.7%; Score 65.6; DB 8;

Best Local Similarity 88.8%; Pred. No. 2.6e-07;

RESULT 1313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 4;
t Local Similarity 88.8%; Pred. No. 2.6e-07;
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(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 4;
EL Local Similarity 88.8%; Pred. No. 2.6e-07;
                                                       2.7%; Score 65.6; DB 5;
88.8%; Pred. No. 2.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.7%; Score 65.6; DB 3;
88.8%; Pred. No. 2.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAC32285 standard; cDNA; 340 BP.
Human secreted protein 5' EST, SEQ ID NO: 36360.
EP1033401-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
                16-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
                                                                       Best Local Similarity RESULT 1312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity RESULT 1317
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Best Local Similarity
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(GEST ) GENSET.
WO200159063-A2.
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Best Loca-
RESULT 1316
ID AAC322°
DE Humar
PN EPJ
PD 0'
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RESULT 1327
ID ADJ07872 standard; DNA; 347 BP.
DE Human cardiovascular system associated polypeptide-related DNA SegID1260.
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Human cardiovascular system associated polypeptide-related DNA SeqID1261.
US2004005575-A1.
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ID ADJ07877 standard; DNA; 347 BP.
DE Human cardiovascular system associated polypeptide-related DNA SeqID1265.
PN US2004005575-A1.
                                                                      AAS35761 standard; DNA; 347 BP.
Human cardiovascular system antigen genomic DNA SEQ ID No 1261
WO200155321-A2.
                                                                                                                                                                                                                                                               ADE46454 standard; DNA; 347 BP. Human carditovascular system related genomic DNA #20.
Human carditovascular system related genomic DNA #20.
27-MAR-2003.
(HUMA) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 10; Length 347; ELocal Similarity 88.8%; Pred. No. 2.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.7%; Score 65.6; DB 13; Length 347; 88.8%; Pred. No. 2.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.7%; Score 65.6; DB 10; Length 347; 88.8%; Pred. No. 2.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.7%; Score 65.6; DB 10; Length 347;
88.8%; Pred. No. 2.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OB-JAN-2004.
(HUMA-) HUMAN GENOME SCI INC.
(ery Match 2.7%; Score 65.6; DB 13; Length 347;
(ery Match 88.8%; Pred. No. 2.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.7%; Score 65.6; DB 13; Length 347;
88.8%; Pred. No. 2.6e-07;
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2.7%; Score 65.6; DB 4; Length 347; 88.8%; Pred. No. 2.6e-07;
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Human ovarian cancer related cDNA clone SEQ ID NO:7821
WO200192581-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADE44459 standard; DNA; 347 BP.
Human cardiovascular system related genomic DNA #25.
US2003059908-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADE46455 standard, DNA, 347 BP.
Human cardiovascular system related genomic DNA #21
US2003059908-A1.
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88.8%; Pred. No. 2.8e-07;
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88.8%; Pred. No. 2.6e-07;
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Human ovarian cancer DNA marker #18630.
WO200170979-A2.
27-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-MAR-2003.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                            02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
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(HUMA-) HUMAN GENOME SCI INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 1328
                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
                 Best Local Similarity RESULT 1321
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Best Local Similarity
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Human prostate expression marker cDNA 49091.
WO200160860-A2.
                                                                                                                    RESULT
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( HELL) HELIX RES INST.

Query Match

2.7%; Score 65.6; DB 4; Length 431;

Act Toral Similarity 88.8%; Pred. No. 3e-07;
                      Length 394;
                                                                                                                                                                                                                                                                                                                                                                                                                   2.7%; Score 65.6; DB 9; Length 426; 88.8%; Pred. No. 2.9e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 2.7%; Score 65.6; DB 6; Length 439;
Local Similarity 88.8%; Pred. No. 3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.7%; Score 65.6; DB 9; Length 474;
88.8%; Pred. No. 3.1e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human ovarian cancer related cDNA clone SEQ ID NO:321
                    2.7%; Score 65.6; DB 5;
88.8%; Pred. No. 2.8e-07;
                                                                                                                                                                           2.7%; Score 65.6; DB 4;
88.8%; Pred. No. 2.8e-07;
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PA (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.

Query Match

2.7%; Score 65.6; DB 5;

BEST Local Similarity 88.8%; Pred. No. 3.1e-07;

RESULT 1336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human cDNA clone (5'-primer) SEQ ID NO:5333.
EP1074617-A2.
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Human prostate expression marker cDNA 45836,
WO200160860-A2.
  (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
                                                                           AAI84018 standard; cDNA; 395 BP.
Human polynucleotide SEQ ID NO 4078.
WO200164835-A2.
                                                                                                                                                                                                                    T 1331
CH17123 standard; cDNA; 426 BP.
Human adult heart cDNA #1437.
US2003073623-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACH47464 standard; cDNA; 458 BP.
Human infant brain cDNA #1527.
US2003073623-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABL77343 standard; cDNA; 439 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACH23761 standard; cDNA; 474 BP.
Human adult ovary cDNA #2141.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (DRWA) DRWANAC R T.
(LABA/) LABAT I.
(STAC/) STACHE-CRAIN B.
(DICK/) DICKSON M C.
(JONE/) JONES L W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LABAT I.
STACHE-CRAIN B.
DICKSON M C.
                                                                                                                                                                                                                                                                                                                                     (LABA/) LABAT I.
(STAC/) STACHE-CRAIN B.
(DICK/) DICKSON M C.
(JONE/) JONES L W.
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                                                                                                                                                                                            Best Local Similarity
                                                                                                                                                                                                                                                                                                                (DRMA/) DRMANAC R T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DRMANAC R T.
                                Best Local Similarity
RESULT 1330
                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
RESULT 1335
                                                                                                                                  07-SEP-2001.
(HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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  PA (MILL-) N
Query Match
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(MILL-) MILLENNIUM PHARM INC.
(MILL-) MILLENNIUM PHARM INC.
2.7%; Score 65.6; DB 11; Length 503;
ery Match 2.7%; Pred. No. 3.2e-07;
                                                                                                                                                                                                                                                                                                  Length 481;
                                      Length 475;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABL77725 standard; cDNA; 495 BP.
Human ovarian cancer related cDNA clone SEQ ID NO:703
WO200192521-A2.
06-DEC-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAL24230 standard; cDNA; 515 BP.
Human breast cancer expressed polynucleotide 16687.
WO200151628-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABN65221 standard; cDNA; 525 BP.
Human cancer related polynucleotide SEQ ID NO 5188.
WOS00214500-A2.
21-FRB-2002.
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Human breast cancer expressed polynucleotide 7841.
WO200151628-A2.
               (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
ry Match
t Local Similarity 88.8%; Pred. No. 3.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                23-AUG-2001.
(MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
2.7%; Score 65.6; DB 5;
tt Local Similarity 88.8%; Pred. No. 3.2e-07;
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ry Match
t Local Similarity 88.8%; Pred. No. 3.3e-07;
                                                                                                                                                                                                                                                                                               2.7%; Score 65.6; DB 9;
88.8%; Pred. No. 3.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
cy Match
2.7%; Score 65.6; DB 5;
Local Similarity 88.8%; Pred. No. 3.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.7%; Score 65.6; DB 4;
88.8%; Pred. No. 3.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.7%; Score 65.6; DB 6;
88.8%; Pred. No. 3.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                Human prostate expression marker cDNA 38038.
WO200160860-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human prostate expression marker cDNA 50162.
WO200160860-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Breast cancer related marker, seq id 13321
US2003099974-A1.
                                                                                                                                                                                                                                                                                                                                        RESULT 1339
ID ABV38047 standard; cDNA; 485 BP.
                                                                                                     ACH23575 standard; cDNA; 481 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACN92171 standard, DNA, 503 BP.
                                                                                                                       Human adult ovary cDNA #1955
US2003073623-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABV50171 standard; cDNA; 494
                                                                                                                                                                                   (DRWA) DRWANAC R T.
(LABA/) LABAT I.
(STAC/) STACHE-CRAIN B.
(DICK/) DICKSON M C.
(JONE/) JONES L W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                    Local Similarity
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23-AUG-2001
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                                        Query Match
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26

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AAH13412 standard;
                                                 WO200155320-A2.
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Best Local Si
RESULT 1363
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:35473.
WO200157182-A2.
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PA (HUMA-) HUMAN GENOME SCI INC.

PA (HUMA-) HUMAN GENOME SCI INC.

Query Match

Best Local Similarity 88.8%; Pred. No. 3.3e-07;

RESULT 1349

ID AAH13272 standard; CDNA; 539 BP.

DE HUMAN CONA CLONE (3'-primer) SEQ ID NO:10107.

PN EP1074617-A2.

PD 07-FEB-2001.

PA (HELI-) HELIX RES INST.

QUERY Match

QUERY MAtch

QUERY MAtch

DAM: 13.9e-07;

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Human reproductive system related antigen DNA SEQ ID NO: 5695.
vuery Match
2.7%; Score 65.6; DB 6; Length 525;
Beet Local Similarity 88.8%; Pred. No. 3.3e-07;
RESULT 1346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 65.6; DB 9; Length 526; Pred. No. 3.3e-07;
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PN EF1074617-A2.

PD 07-FEB-2001.

PA (HELI-) HELIX RES INST.

Query Match
Best Local Similarity 88.8%; Pred. No. 3.4e-07;

RESUL 1351

ID ABV57130 standard; cDNA; 561 BP.

DE Human prostate expression marker cDNA 57121.

PN W2020160860-A2.

PD 23-AUG-2001.

PA (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
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AAH69839 standard, cDNA, 567 BP.
Human cervical cancer marker nucleic acid 1113.
WO200142467-A2.
14-JUN-2001.
(MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
ery Match
2.7%; Score 65.6; DB 4;
et Local Similarity 88.8%; Pred. No. 3.4e-07;
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PA (IMAR-) IMARX PHARM CORP.
Query Match
Best Local Similarity 88.8%; Pred. No. 3.4e-07;
RESULT 1353
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(MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
2.7%; Score 65.6; DB 5;
2.7%; Score 65.6; DB 
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(MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
2.7%; Score 65.6; DB 4;
ery Match
- form! Similarity 88.8%; Pred. No. 3.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human cervical cancer marker nucleic acid 2538.
WO200142467-A2.
                                                                                                                                                                                                 ACH45689 standard; cDNA; 526 BP.
Human foetal brain cDNA #6414.
US2003073623-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 88.8%; Pr
RESULT 1352
ID AAX77064 standard; DNA; 566 BP.
DE Egr gene.
PN W09925385-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.7%;
                                                                                                                                                                                                                                                                                                                                                                 17-APR-2003.
(DRMA/) DRWANAC R T.
(LABA/) LABAT I.
(STAC/) STACHE-CRAIN B.
(DICK/) DICKSON M C.
(JONE/) JONES L W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity RESULT 1347
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Best Local Si
RESULT 1348
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AAKR2986 standard; DNA; 584 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:37798.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAK66436 standard; DNA; 606 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:21248.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAK77289 standard; DNA; 605 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:33101.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                    ABL97354 standard; DNA; 569 BP.
Human testicular antigen encoding DNA fragment SEQ ID NO: 2006.
WO200155317-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 642;
                                           2.7%; Score 65.6; DB 4; Length 569; 88.8%; Pred. No. 3.4e-07;
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                                                                                                                                                                                                                                  Length 569;
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                                                                                                                                                                                                                                                                                                                                                                                                                           Length 569,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 584
                                                                                                               AAH13412 standard; cDNA; 569 BP.

Human cDNA clone (3'-primer) SEQ ID NO:10247.

DOT-FEB-2001.

(HELI-) HELIX RES INST.

Query Match

Best Local Similarity 88.8%; Pred. No. 3.4e-07;
                                                                                                                                                                                                                                                                                                                                                                O2-AUG-2001.
(HUMA) HUMAN GENOME SCI INC.
(HUMA) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 4;
ery Match
2.7%; Pred. No. 3.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MCACCE 09-MG-2001.
(99-MG-2001.
(HUMA-) HUMAN GENOME SCI INC.
Lery Match 2.7%; Score 65.6; DB 4;
Lery Match 88.8%; Pred. No. 3.5e-07;
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(MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.

Query Match
2.7%; Score 65.6; DB 5;

Best Local Similarity 88.8%; Pred. No. 3.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2.7%; Score 65.6; DB 4;
88.8%; Pred. No. 3.7e-07;
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88.8%; Pred. No. 3.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.7%; Score 65.6; DB 4;
88.8%; Pred. No. 3.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MILL-) MILLENNIUM PREDICTIVE MEDICINE INC.
2.7%; Score 65.6; DB 5;
t Local Similarity 88.8%; Pred. No. 3.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAS44682 standard; DNA; 642 BP.
Human full-length polynucleotide sequence #107.
WO200164834-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human prostate expression marker cDNA 50731.
WO200160860-A2.
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Human cDNA clone (5'-primer) SEQ ID NO:5266.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABV08133 standard; cDNA; 586 BP.
Human prostate expression marker cDNA 8124.
WO200160860-A2.
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02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
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Best Local Similarity
RESULT 1358
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Best Local Similarity
                                                                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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(HYSE-) HYSEQ INC.
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PD 30-JAN-2003.
PA (MITS-) MITSUBISHI PHARMA CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FENG P.
LAFLEUR D W.
MOORE P A.
                                                                                                                                          Best Local Similarity RESULT 1373
                                                                                                                                                                                                                                                                                                                            Best Local Similarity RESULT 1374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CARTER K C.
ENDRESS G A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
RESULT 1375
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(RUBE/) RUBEN S M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (WEIY/) WEI Y.
(FANP/) FAN P.
(ROSE/) ROSEN C A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABZ80046 standard;
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KYAW H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200166719-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZENG Z.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (FENG/)
(LAFL/)
(MOOR/)
(SHIY/)
                                                                                                                                                                                                                                                                                                                 Query
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAD31368 standard; DNA; 819 BP.
Amplicon 3, Gscn12.668.05 of 92Kb gene fragment of human chromosome 17.
WO200210455-A2.
                                                                                                                                                                                                                                                                                       AAK67739 standard; DNA; 713 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:22551.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAK65399 standard; DNA; 815 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:20211.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immune/haematopoietic antigen genomic sequence SEQ ID NO:22550.
                                                                                                                                                                                                                                                                                                                                                                                           Length 713;

LucyL57183-A2.

PA (HUMA-) HUMAN GENOME SCI INC.

Query Match
PEEL Local Similarity 88.8%; Pred. No. 3.9e-07;

RESULT 1367

ID AAH04073 standard; CDNA; 786 BP.

DE Human CDNA clone (5'-primer) SP.

Phy (HUMA-) Processed Standard; CDNA; 786 BP.

Phy EP1074617-A2.

Phy (Human CDNA clone (5'-primer) SP.

                                                                                                                                                                          29-MAY-2003.
(MILL-) MILLENNIUM PHARM INC.
bry Match
Bry Match 2.7%; Score 65.6; DB 11; Length 707;
Pt Local Similarity 88.8%; Pred. No. 3.9e-07;
                                           Length 685;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-AUG-2001.
(HUWAN) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 4; Length 815;
st Local Similarity 88.8%; Pred. No. 4.2e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 786;
                                                                                                                                                                                                                                                    ra (HUMA-) HUMAN GENOME SCI INC.

Query March
Best Local Similarity 88.8%; Pred. No. 3.9e-07;
RESULT 1366
ID AAK67738 standard; DNA; 723 BP.
DB Human immune/haematopoietic PP N W0200157187-A2.
PA (HUMA-) HUMAN GENOME SCI INC.

2.7%; Score 65.6; DB 4; Len.

Best Local Similarity 88.8%; Pred. No. 3.9e-07;

RESULT 1366
ID AAK67738 standard; DNA; 723 BP.

PP (HUMAN 1 MM W0200157187-A2.)

PP (HUMAN 1 MM W0200157187-A2.)
07-FEB-2001.
(HELL-) HELIX RES INST.
2.7%; Score 65.6; DB 4;
st Local Similarity 88.8%; Pred. No. 3.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.7%; Score 65.6; DB 6;
Local Similarity 88.8%; Pred. No. 4.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PD 07-FEB-2001.

PA (HELL-) HELIX RES INST.

Query Match 2.7%; Score 65.6; DB 4;

Best Local Similarity 88.8%; Pred. No. 4.1e-07;

RESULT 1368
                                                                                                                               Breast cancer related marker, seq id 6529 US2003099974-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ACD07899 standard; DNA; 821 BP.
DNA encoding human EDA1-II exon 5.
US2003023991-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABL51013 standard, DNA; 821 BP.
Human EDA1-II exon 5 SEQ ID NO:7.
US6355782-B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (BAYU ) BAYLOR COLLEGE MEDICINE. (UYOR-) UNIV OREGON HEALTH SCI.
                                                                                                          ACN85379 standard; DNA; 707 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CELL-) CELLTECH R & D INC. (STRA/) STRAEHLING HAMPTON K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ZONA/) ZONANA J.
(FERG/) FERGUSON B M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (HEAD/) HEADON D. (OVER/) OVERBEEK P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-JAN-2003
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                                                                                                                                                                                                                      Query Match
Best Local Sir
RESULT 1365
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                                              Query Match
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RESULT 1370
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AAK87123 standard; DNA; 898 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:41935.
WO200157182-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            vuery Match 2.7%; Score 65.6; DB 10; Length 824; Best Local Similarity 88.8%; Pred. No. 4.2e-07; RESULT 1376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2.7%; Score 65.6; DB 10; Length 824; 88.8%; Pred. No. 4.2e-07;
AAXX0483 standard; DNA; 824 BP.

Human secreted protein gene 72.

WOS906423-A1.

11-FEB-1999.

(HUMA-) HUMAN GENOME SCI INC.

2.7%; Score 65.6; DB 2; Length 824; ery Match
:st Local Similarity 88.8%; Pred. No. 4.2e-07;
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ry Match 2.7%; Score 65.6; DB 6; Length 824;
t Local Similarity 88.8%; Pred. No. 4.2e-07;
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PA (HISM ) HISAMITSU PHARM CO LTD.
Query Match
2.7%; Score 65.6; DB 4; Length 834;
Best Local Similarity 88.8%; Pred. No. 4.3e-07;
RESULT 1377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 855;
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Human neuroblastoma expressed polynucleotide SEQ ID NO 680.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              y Match 2.7%; Score 65.6; DB 10;
Local Similarity 88.8%; Pred. No. 4.4e-07;
                                                                                                                                                                                                                                           ABQS4154 standard; cDNA; 824 BP.
Human ovarian antigen HRDEL61 cDNA, SEQ ID NO:34
WO200200677-A1.
03-GAN-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADD90267 standard; cDNa; 824 BP.
Novel human secreted protein cDNA seq id 82.
US2003199683-A1.
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Human prostate expression marker cDNA 16038.
WO200160860-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADG90086 standard; cDNA; 824 BP.
Human cDNA from secreted protein gene 72.
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Secreted protein-encoding gene 122 cDNA clone HNHOD46, SEQ ID NO:132.
WO200276488-A1.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:30852.
WO200157182-A2.
09-AUG-2001.
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                                        Length 1314;
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RESULT 1391
ID ACC50566 standard; cDNA; 1355 BP.
DE Human secreted protein coding sequence, SEQ ID 233.
PN W020029510-A2.
PD 28-NOV-2002.
PA (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HUMA-) HUMAN GENOME SCI INC.

ry Match 2.7%; Score 65.6; DB 10;

Local Similarity 88.8%; Pred. No. 5.6e-07;
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(HUMA-) HUMAN GENOME SCI INC.
(ery Match 2.7%; Score 65.6; DB 8;
ery Match 2.7%; Pred. No. 5.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-7AN-2003.
(HUWA-) HUMAN GENOME SCI INC.
(HUWA-) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 9;
Lery Match 2.7%; Pred. No. 5.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2.7%; Score 65.6; DB 8;
88.8%; Pred. No. 5.6e-07;
                                        2.7%; Score 65.6; DB 5;
88.8%; Pred. No. 5.5e-07;
                                                                                                                                                                                                                     2.7%; Score 65.6; DB 3;
88.8%; Pred. No. 5.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.7%; Score 65.6; DB 8;
88.8%; Pred. No. 5.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACT73629 standard; DNA; 1355 BP.

Human secreted protein-related DNA - SEQ ID 262.
W02003038063-A2.
W0-WAY-2003.
(HUMA-) HUMAN GENOME SCI INC.
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ID ADD37644 standard; cDNA; 1355 BP.

DE Human secreted protein encoding sequence #126.

PP WO202029555-A2.

PD 14-NOV-2002.
                                                                                                      AACC9568 standard; DNA; 1355 BP.
Human secreted protein gene 57 clone HNHOD46.
WO200061623-A1.
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Gene encoding human secreted protein #385.
WO2002102994-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human secreted protein cDNA #SEQ ID 177 WO2003004622-A2.
                                                                                                                                                                                                                                                                     ADA40020 standard; cDNA; 1355 BP.
Human secreted protein encoding cDNA.
WQ2002102993-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABZ71311 standard; cDNA; 1355 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADB91231 standard; cDNA; 1355 BP
               PA (HUMA-) HUMAN GENOME SCI INC.
Query Match
Best Local Similarity 88.8%;
RESULT 1389
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-OCT-2002.
(HUMA-) HUMAN GENOME SCI INC.
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                                                                                                                                                                                                 (HUMA-) HUMAN GENOME SCI INC
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Best Local Similarity
RESULT 1392
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Best Local Similarity
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Best Local Similarity
16-AUG-2001
                                                                                                                                                                          19-OCT-2000
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                                                                                                    AAX72553 standard; DNA; 898 BP.

Human immune/haematopoietic antigen genomic sequence SEQ ID NO:27365.

W0200157182-A2.

W0200157182-A2.

(90-Aug-2001.

(HUMA-) HUMAN GENOME SCI INC.

ery Match

2.7%; Score 65.6; DB 4; Length 898;

et Local Similarity 88.8%; Pred. No. 4.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immune/haematopoietic antigen genomic sequence SEQ ID NO:37065.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAK66435 standard; DNA; 1100 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:21247.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF88859 standard; cDNA; 1268 BP.
Phosphoenolpyruvate dependent saccharophosphotransferase 9.46
CN1352250-A.
                                                                                                                                                                                                                                                                                                                                       PD 02-JAN-2003.

PA (NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.

PA (ADSC-) CENT ADVANCED SCI & TECHNOLOGY INCUBATIO.

Query Match 2.7%; Score 65.6; DB 10; Length 931;

Best Local Similarity 88.8%; Pred. No. 4.5e-07;

RESULT 1382
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 2.7%; Score 65.6; DB 3; Length 1092; Best Local Similarity 88.8%; Pred. No. 5e-07; RESULT 1386
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ery Match 2.7%; Score 65.6; DB 4; Length 1100;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 1024;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 1268
                                        Length 898;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABA20555 standard; DNA; 1064 BP.
Human nervous system related polynucleotide SEQ ID NO 12886.
WQ200159063-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NO:41
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ABAIGILI standard, DNA; 1314 BP.
Human nervous system related polynucleotide SEQ ID NO 8442.
WO200159063-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ID . AAC81263 standard; DNA; 1024 BP.

DE Human tyrosine phosphatase HD-PTP gene exon 2, SEQ ID w0200063392-A1.

PD 26-OCT-2000.

PA (KYOW ) KYOWA HAKKO KOUND W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PD 05-JUN-2002.
PA (BODE-) BODE GENE DEV CO LTD SHANGHAI.

QUETY MATCH

BEST Local Similarity 88.8%; Pred. No. 5.4e-07;

RESULT 1388

ID ABA16111 standard; DNA; 1314 PP

DE Human nervous sver-
PN WOODS.
                         vuery Match 2.7%; Score 65.6; DB 4; Best Local Similarity 88.8%; Pred. No. 4.4e-07; RESULT 1380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PD 26-OCT-2000.

PA (KYOW) KYOWA HAKKO KOGYO KK.

Query Match 2.7%; Score 65.6; DB 3;
Best Local Similarity 88.8%; Pred. No. 4.8e-07;

RESULT 1384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
(ery Match 2.7%; Score 65.6; DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human secreted protein gene 6 clone HSJAN83. WO200058358-A1. 05-OCT-2000. (HUMA-) HUWAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA; 1002 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 88.8%; Pre
RESULT 1385
ID AAC79002 standard; DNA; 1092 BP.
                                                                                                                                                                                                                                                                              ADC87114 standard; DNA; 931 BP.
Human GPCR gene SEQ ID NO:1567.
EP1270724-A2.
                  A (HUMA-) HUMAN GENOME SCI INC.
Query Match 2.7%;
                                                                                                                                                                                                                                       Best Local Similarity RESULT 1381
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity RESULT 1383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAK82253 standard;
09-AUG-2001
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Best Local Si
RESULT 1387
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AAS39939 standard; DNA; 1708 BP.
Genomic sequence #358 encoding human colon associated polypeptide.
WQ200155302-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAK90383 standard; DNA; 1708 BP.
Human digestive system antigen genomic sequence SEQ ID NO: 3959.
WO200155314-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human digestive system antigen genomic sequence SEQ ID NO: 4094 WO200155314-A2.
         2.7%; Score 65.6; DB 10; Length 1604;
88.8%; Pred. No. 6.1e-07;
                                                                                                                                                                                   Length 1636;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 1708;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 1701;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.7%; Score 65.6; DB 9; Length 1708; 88.8%; Pred. No. 6.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 1714;
                                                                                                                                                                                                                                                                                                                                                             Length 1643
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 1643;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 1708,
                                                                                                                                                                                                                                               ABA19996 standard, DNA, 1643 BP.
Human nervous system related polynucleotide SEQ ID NO 12327.
WO200159063-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                            ABA19995 standard, DNA, 1643 BP.
Human nervous system related polynucleotide SEQ ID NO 12326.
WO200159063-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADB122899 standard; DNA; 1708 BP.
Human novel colon related polypeptide DNA SEQ ID NO 836.
US2003050231-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human soft tissue sarcoma-upregulated DNA - SEQ ID 8241
WO2004048938-A2.
                                                                                              23
                                                                  ADC35139 standard; cDNA; 1636 BP.
Human breast cancer antigen polynucleotide seg id
US2003108888-A1.
                                                                                                                                                       (LUDW-) LUDWIG INST CANCER RES.

2.7%; Score 65.6; DB 10;
t Local Similarity 88.8%; Pred. No. 6.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PD 07-FEB-2001.
PA (HELI-) HELIX RES INST.
QUETY MATCh
Best Local Similarity 88.8%; Pred. No. 6.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2.7%; Score 65.6; DB 5; 88.8%; Pred. No. 6.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 4;
t Local Similarity 88.8%; Pred. No. 6.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
(HTMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 65.6; DB 5;
ery Match 6.38-07;
                                                                                                                                                                                                                                                                                                                                      (HUMA-) HUMAN GENOME SCI INC.

2.7%; Score 65.6; DB 5;

L Local Similarity 88.8%; Pred. No. 6.2e-07;
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88.8%; Pred. No. 6.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 1410
ID AAH17199 standard; cDNA; 1701 BP.
E Human cDNA sequence SEQ ID NO:16569.
PN EP1074617-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAK90518 standard; DNA; 1714 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADQ25421 standard; DNA; 1714 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-AUG-2001. (HUMA-) HUMAN GENOME SCI INC.
                       Best Local Similarity RESULT 1407
                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                   Best Local Similarity RESULT 1408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ROBE/) ROSEN C A.
(RUBE/) RUBEN S M.
(BARA/) BARASH S C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                       12-JUN-2003
                                                                                                                                                                                                                                                                                                                      16-AUG-200]
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                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                Query Match
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         Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AADOSS83 standard; cDNA; 1604 BP.
Human secreted protein-encoding gene 5 cDNA clone HWDAH38, SEQ ID NO:15.
WO200134627-A1.
                                                                                AAK76042 standard; DNA; 1371 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:30854.
WO200157182-A2.
                                                                                                                                                                                                                                                                                           Human immune/haematopoietic antigen genomic sequence SEQ ID NO:30853
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABK83662 standard; cDNA; 1413 BP.
Human cDNA differentially expressed in granulocytic cells #233
WO200228999-A2.
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(HUWA-) HUWAN GENOME SCI INC.
ery Match
ery Match 2.7%; Score 65.6; DB 4; Length 1604;
                             Length 1369;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.7%; Score 65.6; DB 8; Length 1604;
88.8%; Pred. No. 6.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                  Length 1371;
                                                                                                                                                                                                        Length 1371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 1380
                                                                                                                                                                                                                                                                                                                                                                                                                            ABA21255 standard; DNA; 1380 BP.
Human nervous system related polynucleotide SEQ ID NO 13586.
WO200159063-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACC56676 standard; cDNA; 1604 BP.
Human secreted protein coding sequence, SEQ ID 343.
WO200295010-A2.
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11-APR-2002.

(GENE J. GENE LOGIC INC.

2.7%; Score 65.6; DB 6;

ery Match

2.7%; Score 65.6; DB 6;

ery Match
(HUWA-) HUWAN GENOME SCI INC.
2.7%; Score 65.6; DB 4;
it Local Similarity 88.8%; Pred. No. 5.6e-07;
                                                                                                                                                                                                                                                                                                                                      09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 4;
st Local Similarity 88.8%; Pred. No. 5.6e-07;
                                                                                                                                                                              (HUMA-) HUMAN GENOME SCI INC.
ry Match
L Local Similarity 88.8%; Pred. No. 5.6e-07;
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2.7%; Score 65.6; DB 8;
t Local Similarity 88.8%; Pred. No. 6.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2.7%; Score 65.6; DB 5;
88.8%; Pred. No. 5.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-DEC-2001.
(BIOW-) BIOWINDOW GENE DEV INC SHANGHAI.
xry Macch 2.7%; Score 65.6; DB 6;
rry Macch 88.8%; Pred. No. 5.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADAS6401 standard; DNA; 1604 BP.
Gene encoding human secreted protein #580.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADA40239 standard; cDNA; 1604 BP.
Human secreted protein encoding cDNA.
WO2002102993-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human argininase 9-encoding cDNA. WO200198502-Al.
                                                                                                                                                                                                                                                    T 1399
AAK76041 standard; DNA; 1371 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-AUG-2001. (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (HUMA-) HUMAN GENOME SCI INC.
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                       Query Match
Best Local Similarity
RESULT 1398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        local Similarity
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Best Local Similarity
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                                                                                                                                                              09-AUG-200
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                                                                                                                                                                                                     Query Match
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Best RESULT

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Human secreted protein-encoding gene 12 cDNA clone HNHEI85, SEQ ID NO:22. WO200132676-A1.
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Human immune/haematopoietic antigen genomic sequence SEQ ID NO:25606.
WO200157182-A2.
                                                                                                                                                                  PA (HELL-) HELIX RES INST.
PA (REAS-) RES ASSOC BIOTECHNOLOGY.
Query Match 2.7%; Score 65.6; DB 10; Length 1939;
Best Local Similarity 88.8%; Pred, No. 6.8e-07;
RESULT 1426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.7%; Score 65.6; DB 12; Length 2175; 88.8%; Pred. No. 7.3e-07;
2.7%; Score 65.6; DB 10; Length 1930;
88.8%; Pred. No. 6.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PA (HELI-) HELIX RES INST.
Query Match
2.7%; Score 65.6; DB 4; Length 2012;
Best Local Similarity 88.8%; Pred. No. 6.9e-07;
RESULT 1429
                                                                                                                                                                                                                                                                                                                                                                                                                    Length 1962;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ouery Match 2.7%; Score 65.6; DB 4; Length 1997; Best Local Similarity 88.8%; Pred. No. 6.9e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 2043;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA81645 standard; DNA; 2140 BP.
N. meningitidis partial DNA sequence gnm_192 SEQ ID NO:192.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
(HTMA-) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 4;
lery Match 2.7%; Pred. No. 7.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                          (HELI-) HELIX RES INST.
2.7%; Score 65.6; DB 4;
Best Local Similarity 88.8%; Pred. No. 6.9e-07;
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88.8%; Pred. No. 7.2e-07;
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PA (BIOW-) BIOWINDOW GENE DEV INC SHANGHAI.

Query Match 2.7%; Score 65.6; DB 8;

Beet Local Similarity 88.8%; Pred. No. 7e-07;

RESULT 1430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human zinc finger protein 8.91-encoding cDNA.
                                                                    ADB63149 standard; cDNA; 1939 BP.
Human cDNA encoding clone SPLEN20041810.
                                                                                                                                                                                                                                                                                          AAH13715 standard; cDNA; 1962 BP.
Human cDNA sequence SEQ ID NO:10603.
EP1074617-A2.
O7-FEB-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAH16700 standard; cDNA; 2012 BP.
Human cDNA sequence SEQ ID NO:15865.
EP1074617-A2.
07-FEB-2001.
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Human cDNA sequence SEQ ID NO:15831
EP1074617-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 1427
ID AAD07582 standard; cDNA; 1997 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADQ64339 standard; cDNA; 2175 BP.
Novel human cDNA sequence #1500.
EP1440981-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABZ57045 standard; cDNA; 2043 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (REAS-) RES ASSOC BIOTECHNOLOGY.
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(HUMA-) HUMAN GENOME SCI INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-APR-2000.
(CHIR ) CHIRON CORP.
                      Best Local Similarity RESULT 1425
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity RESULT 1432
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
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                                                                                                                       EP1308459-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-JUL-2004
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  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADR07263 standard; cDNA; 1840 BP. Full length human cDNA useful for treating neurological disease Seq 769.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 1419
ID AAK82508 standard; DNA; 1845 BP.
DE Human immune/haematopoietic antigen genomic sequence SEQ ID NO:37320.
PN WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAK83510 standard; DNA; 1845 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:37322.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAK82209 standard; DNA; 1845 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:37321.
WO200157182-A2.
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Human insulin resistance marker IRM330 (G protein gamma-4) cDNA #2.
WO200298355-A2.
                                                                                                                                                                                                                                                                                                                        #
                                                                                                                                                                                                                                                                                          ADA13372 standard; cDNA; 1734 BP.
Human insulin resistance marker IRM330 (G protein gamma-4) cDNA
WO200298355-A2.
                                                                                                                                                                                                                                                                                                                                                                                                          Ouery Match
Best Local Similarity 88.8%; Pred. No. 6.4e-07;
RESULT 1418
DE RAD7263 standard; CDNA; 1840 BP.
DE Pull length human CDNA useful for treating neurological disease
PD 8P14413-A2.
PD 18-AUG-2004.
PA (REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                  PD 10-UUN-2004.

PA (PROT-) PROTEIN DESIGN LABS INC.

Query Match

2.7%; Score 65.6; DB 12; Length 1714;

Best Local Similarity 88.8%; Pred. No. 6.4e-07;

RESULT 1417
                         2.7%; Score 65.6; DB 12; Length 1714; 88.8%; Pred. No. 6.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2.7%; Score 65.6; DB 12; Length 1863; 88.8%; Pred. No. 6.7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 1840
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88.8%; Pred. No. 6.7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-AUG-2001.
(HUMA.) HUMAN GENOME SCI INC.
12.7%; Score 65.6; DB 4; Length 1845;
st Local Similarity 88.8%; Pred. No. 6.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-AUG-2001.
(HUMA) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 4; Length 1845;
it Local Similarity 89.8%; Pred. No. 6.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 1845,
                                                                                              ADQ23971 standard; DNA; 1714 BP.
Human soft tissue sarcoma-upregulated DNA - SEQ ID 6791
WO2004048938-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AD021477 standard; DNA; 1863 BP.
Human soft tissue sarcoma-upregulated DNA - SEQ ID 6297
WO2004048938-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     D 18-AUG-2004.
A (REAS-) RES ASSOC BIOTECHNOLOGY.
Query Match
2.7%; Score 65.6; DB 13;
Best Local Similarity 88.8%; Pred. No. 6.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PD 09-AUG-2001.

PA (HUMA-) HUMAN GENOME SCI INC.

QUETY MATCh

Best Local Similarity 88.8%; Pred. No. 6.6e-07;

RESULT 1421
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ID AAL51671 standard; cDNA; 1886 BP.

DE Signal peptidase 11-77 coding sequence.

RN1360046-A.

PD 24-JUL-2002.

PA (BODE-) BODE GENE DEV CO LTD SHAMMINAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-JUL-2002.
(BODE-) BODE GENE DEV CO LTD SHANGHAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-JUN-2004.
(PROT-) PROTEIN DESIGN LABS INC.
     (PROT-) PROTEIN DESIGN LABS INC.
                                                                                                                                                                                                                                                                                                                                                                         12-DEC-2002. (CLIN-) CLINGENIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CLIN-) CLINGENIX INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 1424
                                           Best Local Similarity RESULT 1416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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RESULT 1420
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BBBBBB

Length 2604;

.; 8

Length 2629;

87

Query Match

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ADRO7439 standard; cDNA; 2651 BP.
Full length human cDNA useful for treating neurological disease Seq 945.
EP1447413-A2.
                                                                                                                                                                                                                                                                                                                                                                      (HUMA-) HUMAN GENOME SCI INC.

2.7%; Score 65.6; DB 10; Length 2604;

t Local Similarity 88.8%; Pred. No. 8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CHBF) GRUENENTHAL GMBH.
2.7%; Score 65.6; DB 13; Length 2630; t Local Similarity 88.8%; Pred. No. 8.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.7%; Score 65.6; DB 12; Length 2636; 88.8%; Pred. No. 8.1e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.7%; Score 65.6; DB 13; Length 2651; 88.8%; Pred. No. 8.1e-07;
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Novel human secreted and transmembrane protein cDNA SegID 145.
WO2003072035-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ouery Match 2.7%; Score 65.6; DB 10; Length 2642; Best Local Similarity 88.8%; Pred. No. 8.1e-07;
                   2.7%; Score 65.6; DB 8; Length 2604; 88.8%; Pred. No. 8e-07;
                                                                                                                                                                                                                                                                           ABZ67754 standard; DNA; 2604 BP.
Human secreted protein encoding genomic DNA SEQ ID NO 1277.
WO200277186-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAS74215 standard; cDNA; 2629 BP.
DNA encoding novel human diagnostic protein #10019.
WO200175067-A2.
                                                                                          Human secreted protein-related DNA sequence #310. 16-JAN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.7%; Score 65.6; DB 5;
88.8%; Pred. No. 8.1e-07;
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                                                                                                                                                                                                         2.7%; Score 65.6; DB
88.8%; Pred. No. 8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH17727 standard; cDNA; 2642 BP.
Human cDNA sequence SEQ ID NO:17334.
EP1074617-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADA52774 standard; cDNA; 2684 BP.
Human coding sequence, SEQ ID 342.
EP1293569-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AD064717 standard; cDNA; 2636 BP.
Novel human cDNA sequence #1878.
EP1440981-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CDNA; 2630 BP
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(REAS-) RES ASSOC BIOTECHNOLOGY.
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(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                   ADA98717 standard; DNA; 2604 BP
                                                                                                                                                                                   (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-FEB-2001.
(HELL-) HELIX RES INST.
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(GETH ) GENENTECH INC.
  Best Local Similarity RESULT 1443
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (HYSE-) HYSEQ INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human RAB21 cDNA.
WO2004070383-A2.
19-AUG-2004.
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Secreted protein gene 215 genomic fragment HNGAM58, SEQ ID NO:1335.
WO200277013-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 2.7%; Score 65.6; DB 11; Length 2405; Best Local Similarity 88.8%; Pred. No. 7.7e-07; RESULT 1439
Query Match 2.7%; Score 65.6; DB 4; Length 2191;
Best Local Similarity 88.8%; Pred. No. 7.3e-07;
RESULT 1434
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 2.7%; Score 65.6; DB 12; Length 2242; Local Similarity 88.8%; Pred. No. 7.4e-07;
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                                                          ABQ80846 standard; cDNA; 2195 BP.
Tyrosine specific protein phosphatase 23.87 coding sequence
CN1352275-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAC75794 standard; cDNA; 2468 BP.
Human ORFX ORF1349 polynucleotide sequence SEQ ID NO:2697.
WO200058473-A2.
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Lung cancer associated polynucleotide sequence SEQ ID 171
WO200055180-A2.
                                                                                                                                                                                                                                                          ABL41381 standard; cDNA; 2195 BP.
Human protein phosphatase 8.8 encoding cDNA SEQ ID NO
CN1327061-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADR83499 standard; DNA; 2428 BP.
Human VENT-like homeobox 2 DNA, target gene of miRNA.
WO2004076622-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Absolute standard; cDNA; 2332 BP.
Human secretory polynucleotide SPTM SEQ ID NO 286.
94.000 20293876-A2.
                                                                                                                                                                                                                                                                                                                                           19-DEC-2001.
(BODE-) BODE GENE DEV CO LTD SHANGHAI.
2.7%; Score 65.6; DB 6;
et Local Similarity 88.8%; Pred. No. 7.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-SEP-2004.
(NAAD-) NAT INST ADVANCED IND SCI & TECHNOLOGY.
2.7%; Score 65.6; DB 13;
st Local Similarity 88.8%; Pred. No. 7.7e-07;
                                                                                                                                                        05-JUN-2002.
(BODE-) BODE GENE DEV CO LTD SHANGHAI.
2.7%; Score 65.6; DB 6;
et Local Similarity 88.8%; Pred. No. 7.3e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2.7%; Score 65.6; DB 8;
Local Similarity 88.8%; Pred. No. 7.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADM01346 standard; cDNA; 2405 BP.
Human cDNA of the invention SEQ ID NO:31.
EP1347046-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-SEP-2003.
(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CDNA; 2242 BP
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(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADQ64167 standard; cDNA; 2242 BP
Novel human cDNA sequence #1328.
EP1440981-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-OCT-2002.
(INCY-) INCYTE GENOMICS INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-OCT-2000.
(CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
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Length 2642;

Query Match

Length 2914;

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ADR07894 standard; cDNA; 2891 BP.
Full length human cDNA useful for treating neurological disease Seq 1400.
EP1447413-A2.
                                                                                                                                                                                                                                        AAK70730 standard; DNA; 2914 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:25542.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immune/haematopoietic antigen genomic sequence SEQ ID NO:40050.
W0200157182-A2.
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Pred. No. 8.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2.7%; Score 65.6; DB 11; Length 2978; 88.8%; Pred. No. 8.6e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 65.6; DB 11; Length 2981; Pred. No. 8.6e-07;
                                                                                                                                                                                    2.7%; Score 65.6; DB 13; Length 2891;
88.8%; Pred. No. 8.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human purified secretory polynucleotide (sptm), seq id 47.
WO2003062385-A2.
                                                                                                                                                                                                                                                                                        09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
ery Match 2.7%; Score 65.6; DB 4;
ery Match 88.8%; Pred. No. 8.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.7%; Score 65.6; DB 4;
88.8%; Pred. No. 8.7e-07;
                                                              88.8%; Pred. No. 8.5e-07;
                                                                                                                                                                                                                                                                                                                                                                                    ACN88617 standard; DNA; 2978 BP.
Breast cancer related marker, seq id 9767.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACN45127 standard; cDNA; 2981 BP.
Human mRNA sequence hCT5626,
WO2003073826-A2.
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                                                                                                                                                   18-AUG-2004.
(REAS-) RES ASSOC BIOTECHNOLOGY.
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(HUMA-) HUMAN GENOME SCI INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAK85238 standard; DNA; 3015
                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MILL-) MILLENNIUM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-SEP-2003.
(SAGE-) SAGRES DISCOVERY.
BY MATCH 2-7%;
St Local Similarity 88.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INCY-) INCYTE GENOMICS INC.
JONE/) JONES A L.
DAHL/) DAHL C R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LAN R Y.
URASHKA M E.
KRISTNAM S R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CHINN J.
DUFOUR G E.
JACKSON J L.
                                                            Best Local Similarity RESULT 1461
                                                                                                                                                                                      Query Match
Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
RESULT 1467
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                              Best Local Similarity RESULT 1463
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DAHL C R.
GIETZEN D.
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AMSHEY S R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DAM T C.
LIU T F.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CHEN A J.
                                                                                                                                                                                                                                                                                                                                                                                                                       US2003099974-A1.
29-MAY-2003.
                                                                                                                                                                                                                                                                               WO200157182-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1-JUL-2003
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AMSH/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GIET/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MARW/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABK42391 standard; DNA; 2867 BP.
Genomic sequence #290 encoding novel human connective tissue polypeptide.
WO200155343-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADJ08711 standard; DNA; 2687 BP.
Human cardiovascular system associated polypeptide-related DNA SegID2099.
US2004005575-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 1456
ID AD08712 standard, DNA, 2687 BP.
DE Human cardiovascular system associated polypeptide-related DNA SeqID2100.
PN US2004005575-A1.
           Loure 65.6; DB 10; Length 2684;

Loure 65.6; DB 10; Length 2684;

Luman cardiovascular system antigen genomic DNA SEQ ID No 2099.

Ph W0200155321-A2.

PA (HUMA-) HUMAN GENOME SCI INC.

Query Match

Best Local Similarity 88.8%; Pred. No. 8.2e-07;

RESULT 1453

ID AAS36600 standard; DNA; 2687 BP.

DE Human cardiovascular system and accidence and accidence and accidence accidence.

PN W0200155321-A2.

PD 02-AUG-20n1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 2687;
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2.7%; Score 65.6; DB 9; Length 2867;
rry Match
                                                                                                                                                                                                                                                                                                                                    weir match 2.7%; Score 65.6; DB 4; Length 2697; Best Local Similarity 88.8%; Pred. No. 8.2e-07; RESULT_1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 2716;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2.7%; Score 65.6; DB 4; Length 2867;
88.8%; Pred. No. 8.5e-07;
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PA (HUMA-) HUMAN GENOME SCI INC.

Query Match

2.7%; Score 65.6; DB 10; Let

Query Match

2.7%; Pred. No. 8.2e-07;

Rest Local Similarity 88.8%; Pred. No. 8.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADE47294 standard; DNA; 2687 BP.
Human cardiovascular system related genomic DNA #860.
US2003059908-A1.
                                                                                                                                                                                                                                                                                                                                                                                                       ADE47293 standard; DNA; 2687 BP.
Human cardiovascular system related genomic DNA #859.
US2003059908-A1.
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(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 13;
it Local Similarity 88.8%; Pred. No. 8.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-MAR-2003.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 10;
st Local Similarity 88.8%; Pred. No. 8.2e-07;
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ry Match

t Local Similarity 88.8%; Pred. No. 8.2e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human cDNA sequence SEQ ID NO:16401.
EP1074617-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HUMA-) HUMAN GENOME SCI INC.
Jest Local Similarity 6
RESULT 1452
ID AAS36599 stand>
DE Human card*
PN WO2001**
PD 02-*
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Best Local Si
RESULT 1455
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RESULT 1457
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Length 3015,

83

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AAK70074 standard; DNA; 3454 BP
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Best Local Similarity
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RESULT 1479
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity RESULT 1484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABQ81803 standard;
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(GEST ) GENSET.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-MAR-2001.
(GEST ) GENSET.
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                                                                                                                                            AAF71302 standard; DNA; 3108 BP.
Human low adenosine antisense oligonucleotide related sequence #2869.
WO200062736-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAK72154 standard; DNA; 3426 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:26966.
WO200157182-A2.
Human adenosine receptor related polymucleotide 2nd SEQ ID NO:54.

N WO200009525-A2.

D 24-FEB-2000.

A (UFC-) UNIV EAST CAROLINA.

2.7%; Score 65.6; DB 3; Length 3108;

Best Local Similarity 88.8%; Pred. No. 8.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Match 2.7%; Score 65.6; DB 10; Length 3108; Local Similarity 88.8%; Pred. No. 8.8e-07;
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PD (HUMA-) HUMAN GENOME SCI INC.
PA (HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 4; Length 3426;
Query Match 2.7%; Pred. No. 9.3e-07;
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Human nervous system related polynucleotide SEQ ID NO 6690.
WO200159063-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                T 1473
ABA01586 standard; cDNA; 3314 BP.
Human neuroprotein Y 11 encoding cDNA SEQ ID NO:1.
WG200175020-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 2.7%; Score 65.6; DB 10;
Best Local Similarity 88.8%; Pred. No. 9.1e-07;
                                                                                                                                                                                                                                                                              Score 65.6; DB 3;
Pred. No. 8.8e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human pulmonary and inflammatory target DNA #456 WO200285309-A2.
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ry Match
t Local Similarity 88.8%; Pred. No. 8.9e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Match 2.7%; Score 65.6; DB 6; Local Similarity 88.8%; Pred. No. 9.2e-07;
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(HUMA-) HUMAN GENOME SCI INC.

ery Match 2.7%; Score 65.6; DB 5;

ery Match 88.8%; Pred. No. 9.2e-07;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH18730 standard; CDNA; 3164 BP.
Human CDNA sequence SEQ ID NO:19006.
EP1074617-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADA53534 standard; cDNA; 1255 BP.
Human coding sequence, SEQ ID 1102.
EP1293569-A2.
                                                                                                                                                                                                                                                                                                                                            DNA; 3108 BP.
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(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA; 3108 BP.
                                                                                                                                                                                                                                                                            2.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                             (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                       ABZ96996 standard; DNA; 3106
Human nucleic acid sequence.
WO200285308-A2.
                                                                                                                                                                                                                            (UYEC-) UNIV EAST CAROLINA. (NYCE/) NYCE J W.
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Best Local Similarity
RESULT 1475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABD20845 standard;
                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
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RESULT 1472
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ADK61652 standard; DNA; 3693 BP.
Ovarian cancer-related DNA #807 with altered ovarian cancer expression.
WO2003068054-A2.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:24886.
WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human immune/haematopoietic antigen genomic sequence SEQ ID NO:25690. WO200157182-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PA (PROT-) PROTEIN DESIGN LABS INC.
Query Match 2.7%; Score 65.6; DB 12; Length 3654;
Best Local Similarity 88.8%; Pred. No. 9.7e-07;
RESULT 1480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 3572;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (USSH) US DEPT HEALTH & HUMAN SERVICES.
(SLOK ) SLOAN KETTERING INST CANCER RES.

ry Match 2.7%; Score 65.6; DB 10; Length 3693;
t Local Similarity 88.8%; Pred. No. 9.7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2.7%; Score 65.6; DB 6; Length 3809; 88.8%; Pred. No. 9.9e-07;
                                                                                                   Length 3454;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADQ24785 standard; DNA; 3654 BP.
Human soft tissue sarcoma-upregulated DNA - SEQ ID 7605.
WO2004048938-A2.
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ry Match
2.7%; Score 65.6; DB 10;
t Local Similarity 88.8%; Pred. No. 9.7e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 65.6; DB 10;
No. 9.6e-07;
                             PD 09-AUG-2001.
PA (HUMA-) HUMAN GENOME SCI INC.
Query Match
Best Local Similarity 88.8%; Pred. No. 9.4e-07;
RESULT 1477
                                                                                                                                                                                                                                                                                       Query Match 2.7%; Score 65.6; DB 4; Best Local Similarity 88.8%; Pred. No. 9.5e-07; RESULT 1478
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88.8%; Pred. No. 9.9e-07;
                                                                                                                                                                                                                                                                                                                                                                          AD862232 standard; cDNA; 3572 BP.
Human cDNA encoding clone CTONG20084660.
EP1308459-A2.
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3 CDNA SEQ ID NO:2.
                                                                                                                                                                         AAH18248 standard; cDNA; 3543 BP.
Human cDNA sequence SEQ ID NO:18200.
EP1074617-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2.7%; Score
88.8%; Pred.
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Human PG-3 coding sequence.
WO200114550-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-MAY-2003.
(HELI-) HELIX RES INST.
(REAS-) RES ASSOC BIOTECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADF60998 standard; DNA; 3693 BP.
Pain associated human 336 gene.
US2003153525-A1.
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(HUMA-) HUMAN GENOME SCI INC.
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AAKB2280 standard; DNA; 4063 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:37092.
WO200157182-A2.
                                                                                                                                                                                                                                                                      AAK70341 standard; DNA; 4063 BP.
Human immune/haematopoietic antigen genomic sequence SEQ ID NO:25153.
WO200157182-A2.
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(BRIM) BRISTOL-MYERS SQUIBB CO.
(ery Match 2.7%; Score 65.6; DB 13; Length 4314;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.7%; Score 65.6; DB 12; Length 4247; 88.8%; Pred. No. 1.1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2.7%; Score 65.6; DB 12; Length 4314; 88.8%; Pred. No. 1.1e-06;
                     2.7%; Score 65.6; DB 4; Length 3961;
88.8%; Pred. No. 1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 4153;
                                                                                                                                                                                                           Length 4018
                                                                                                                                                                                                                                                                                                                                                                                         Length 4063
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 4063
                                                                                                            Nucleotide sequence of human chemokine receptor CCR6. WO2003014153-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADJ95422 standard; DNA; 4247 BP.
Human nootropic protein coding sequence, SEQ ID 11.
JP2003310264-A.
                                                                                                                                                                                                                                                                                                                              MOLOUG-2001.
(19-MGG-2001.
(HUMA-) HUMAN GENOME SCI INC.
2.7%; Score 65.6; DB 4;
lery Match
2.7%; Pred. No. 1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AUN1440 standard; DNA; 4314 BP.
Human NF-kappaB pathway-associated gene SeqID441.
WO2004065577-A2.
                                                                                                                                                                          (TOPI-) TOPIGEN PHARM INC.
Query Match
Best Local Similarity 88.8%; Pred. No. 1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 2.7%; Score 65.6; DB 4;
Best Local Similarity 88.9%; Pred. No. 1e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PD 18-OCT-2001.

PA (GEMY) GENETICS INST INC.

QUESTY MATCH
2.7%; Score 65.6; DB 6;

Best Local Similarity 88.8%; Pred. No. 1e-06;

RESULT 1498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADP10544 standard; DNA; 4314 BP.
Reference mRNA sequences for marker probe #221.
WO2004042346-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1497
ID ABK34686 standard; CDNA; 4153 BP.
DE Human CDNA for novel secreted protein, SEQ ID
PN WO200177290-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21-MAY-2004.
(EXPR-) EXPRESSION DIAGNOSTICS INC
                                                                                       ABZ68882 standard; cDNA; 4018 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-NOV-2003.
(KAZU-) ZH KAZUSA DNA KENKYUSHO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-AUG-2001.
(HUMA-) HUMAN GENOME SCI INC.
(HUMA-) HUMAN GENOME SCI INC.
                                 Best Local Similarity
RESULT 1494
                                                                                                                                                                                                                      Best Local Similarity
RESULT 1495
                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
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Best Local Similarity
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                       Query Match
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